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(71) Applicant: CHIRON CORPORATION [US/US]; 4560 Horton Street, Emeryville, CA 95608 (US).

(72) Inventors: ZUKERMANN, Ronald; 1126 Keeler Avenue, Berkeley, CA 94708 (US). DUBOIS-STRINGFELLOW, Nathalie: 1008 The Alameda, Berkeley, CA 94707 (US). DWARKI, Varavani; 1177 Old Alameda Pt., Alameda, CA 94502 (US). INNIS, Michael, A.; 315 Constance Place, Moraga, CA 94556 (US). MURPHY, John, E., 49 Hourbord Court, Oakland, CA 94618 (US). COHEN, Fred; Chiron Corporation, Intellectual Property - R440, P.O. Box 8097, Emeryville, CA 94662-8097 (US). TETSUO, Uno; 480 Warren Drive #530, San Francisco, CA 94131 (US).

(74) Agents: FUJITA, Sharon, M. et al.; Chiron Corporation, Intellectual Property - R440, P.O. Box 8097, Emeryville, CA 94662-8097 (US).

(54) Title: COMPOSITIONS AND METHODS FOR POLYNUCLEOTIDE DELIVERY

(57) Abstract

This invention relates to compositions and methods for increasing the uptake of polynucleotides into cells. Specifically, the invention relates to vectors, targeting ligands, and polycationic agents. The polycationic agents are capable of (1) increasing the frequency of uptake of polynucleotides into a cell, (2) condensing polynucleotides; and (3) inhibiting serum and/or nuclease degradation of polynucleotides.

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Compositions and Methods for Polynucleotide Delivery

Description

Field of the Invention

This invention relates to compositions and methods for increasing the uptake of polynucleotides into cells. Specifically, the invention relates to vectors, targeting ligands, and polycationic agents. The polycationic agents are capable of (1) increasing the frequency of uptake of polynucleotides into a cell, (2) condensing polynucleotides; and (3) inhibiting serum and/or nuclease degradation of polynucleotides.

10 Background of the Invention

Polycations, such as polylysine, have been used to facilitate delivery of nucleic acids to cell interior. Both *in vitro* and *in vivo* applications have taken advantage of this property. See, for example, Gao et al., 1996, Biochem. 35:1027-1036.

Polynucleotides, typically DNA, may be taken into a cell by a receptormediated endocytosis pathway, a cellular mechanism which internalizes specific
macromolecules. In general, complexes designed to be delivered in this fashion
contain nucleic acids encoding the gene of interest and a polycationic agent,
which acts as a DNA-binding carrier and both neutralizes the charge on the
nucleic acids and condenses it.

Condensation facilitates entry of nucleic acids into cell vesicle systems by simulating a macromolecular structure. For example, polylysine condenses DNA into a toroid or doughnut-like structure. (Wagner et al., 1991, Proc. Natl. Acad. Sci. 88:4255-4259).

Polycations previously utilized for nucleic acid delivery to ceil interiors include polylysine, protamines, histones, spermine, spermidine, polyornithine, polyargnine, and putrescine.

All publications mentioned herein are incorporated herein by reference for the purpose of disclosing and describing features of the invention for which the publications are cited in connection with.

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Summary of the Invention

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An embodiment of the invention is a vector for expression of polypeptides. The vector of the instant invention comprises: (i) an Epstein Barr Virus (EBV) origin of replication; (ii) a polynucleotide encoding an EBV origin binding protein; (iii) an enhancer; (iv) a promoter; and (v) a terminator. Polynucleotides encoding a desired polypeptide, such as erythropoietin or leptin can be inserted into the vector. Also, ribozyme and antisense polynucleotides can also be inserted into the vector.

One embodiment of the invention is a composition capable of targeting a polynucleotide to a specific cell type. The composition comprises: (i) a lipoprotein; (ii) a polynucleotide binding molecule; and (iii) a polynucleotide.

Another embodiment of the invention is a method of increasing the frequency of uptake of polynucleotides into a cell by contacting a cell with a composition comprising: (i) a lipoprotein, (ii) a polynucleotide binding molecule; and (iii) a polynucleotide.

Yet another embodiment of the invention is a method of increasing the frequency of uptake of polynucleotides into a specific cell type by contacting a population of cells with a composition comprising (i) a lipoprotein, (ii) a polynucleotide binding molecule; and (iii) a polynucleotide.

One embodiment of the invention is a polycationic agent exhibiting a net positive electrical charge at physiological pH with the following formula:

$$Ta - \begin{bmatrix} -\frac{R_1}{N} & \frac{R_2}{N} & 0 \\ -\frac{C}{N} - \frac{C}{C} - C - \end{bmatrix}_{n} - Tc$$

where Ta and Tc are terminating groups. A preferred subset of these compounds is the set where R_2 is hydrogen. Even more preferred are polymers comprising at least one unnatural amino acid. Also preferred are polymers where R_2 and R_3 are hydrogen and R_1 is not hydrogen, also referred to as poly N-substituted glycines or "poly NSGs."

Another embodiment is a neutral polymer exhibiting no net positive or negative electrical charge at physiological pH with the following formula:

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BNon--

$$Ta - \begin{bmatrix} -\frac{R_1}{N} & \frac{R_2}{N} & 0 \\ -\frac{N}{N} - \frac{C}{C} - \frac{C}{C} - \frac{C}{N} \end{bmatrix}_n - Tc$$

where Ta and Tc are terminating groups. A preferred subset of these compounds is the set where R_2 is hydrogen. Even more preferred are polymers comprising at least one unnatural amino acid. Also preferred are polymers where R_2 and R_3 are hydrogen and R_4 is not hydrogen, also referred to as poly N-substituted glycines or "poly NSGs."

The instant polycationic agents and neutral polymers are capable of neutralizing the electrical charge of nucleic acids. A subset of these compounds are capable of (1) condensing the structure of polynucleotides and/ or (2) protecting polynucleotides from serum and/or nuclease degradation.

Yet another embodiment of the invention are polycationic agents and neutral polymers that (1) target binding of nucleic acids to cell surfaces, (2) trigger cell membrane destabilization; (3) exhibit endosome buffering capacity; (4) trigger endocytosis; (5) help trigger the release of polynucleotide/lipid complexes from endosomes or (6) nuclear tropism.

Another embodiment of the invention is a composition comprising a polynucleotide of interest and an effective amount of the polycationic agent to neutralize the charge of the polynucleotide. Optionally, the composition includes a ligand which directs the complex to particular cells expressing a ligand-binding partner, and/or an endosomolytic agent, which serves to cause disruption of the endosome containing the complex.

Another embodiment of the invention is a method of condensing nucleic acids by providing an effective amount of the polycationic agents or neutral polymers of the invention and contacting the agent with the desired polynucleotides.

Also an embodiment of the invention is a method of inhibiting serum and/or nuclease degradation of nucleic acids by providing an effective amount of the the polycationic agents or neutral polymers of the inventions and contacting the agent with the desired nucleic acids.

Brief Description of the Drawings

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Figure 1 is a schematic of a two-step monomer assembly reaction scheme.

Figure 2 is a schematic of a three-step monomer assembly reaction scheme.

Figure 3 is a plasmid map of vector pCMVKmlTR-EPI.

Figure 4 is a plasmid map of vector CMVkm2.

Figure 5 is a plasmid map of vector pCMV-KM-cmEPO.

Figure 6 is a plasmid map of vector CMVKmLeptinWt.

Figure 7A illustrates transfection efficiencies for a diverse set of polycationic agents. The polycatonic agents were formulated with DNA at a 2:1, + to - charge ratio and added to either HT1080 (solid bar) or COS (stippled bar) in the presence of 10% serum. Luciferase activity was analyzed 48 hours post-transfection. Total cell protein was measured using a Pierce BCA assay and luciferase activity was normalized against total cell protein.

Figure 7B illustrates the effect of oligomer length on transfection efficiency for polycationic agents having different numbers of the same repeating trimer motif. For both A and B each data point represents the average of 2 experiments.

Figure 8(A-C) shows RZ145-1 peptoid-mediated transfection and transfection mediated by commercially available cationic liposome preparations. RZ145-1 or the indicated lipid was formulated and added to cells in the presence (solid bar) or absence (stippled bar) of 10% serum. Luciferase and total cell protein activity were measured 48 hours after initial transfection. Cells lines are (Figure 8A) 293 human embryonic kidney cells, (Figure 8B) HT1080 human fibrosarcoma cells, and (Figure 8C) NIH03T3 mouse fibroblast cells. Each data point represents the average + strandard error of the mean of three transfections.

Figure 9 illustrates the effect of chloroquine on transfection with RZ145-1 in (A) 293 cells and (B) HT1080 cells. The cells were transfected in the presence (black bar) or absence (stippled bar) of 100 uM chloroquine. Cells were lysed 48 hours post transfection and luciferase activity and total protein content were measured.

Detailed Description

Definitions

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"Lipoproteins" refers to polypeptides that are associated non-covalently with lipids in the bloodstream and are capable of binding to cellular receptors. Preferably, lipoproteins are those involved with transport and storage of lipids. Such proteins include, for example, chylomicrons, low density lipoprotein (LDL), very low density lipoprotein (VLDL), intermediate density lipoprotein (IDL), and high density lipoprotein (HDL). Also, included in the term are mutants, fragments, or fusions of the naturally occurring lipoproteins. Also, modifications of naturally occurring lipoproteins can be used, such as acetylated LDL.

Mutants, fragments, fusions, or modifications of the naturally occurring lipoproteins are amino acid sequences that exhibit substantial sequence identity to naturally occurring lipoproteins or a fragment thereof. These polypeptides will retain more than about 80% amino acid identity; more typically, more than about 85%; even more typically, at least 90%. Preferably, these polypeptides will exhibit more than about 92% amino acid sequence identity with naturally occurring lipoproteins or fragment thereof; more preferably, more than about 94%; even more preferably, more than about 96%; even more preferably, more than about 98%; even more preferably, more than about 99%. All of these polypeptides exhibit receptor binding properties of naturally occurring lipoproteins. Usually, such polypeptides exhibit at least about 20% receptor binding of naturally occurring lipoproteins. More typically, the polypeptides exhibit at least about 40%, even more typically the polypeptides exhibit at least about 60%; even more typically, at least about 70%; even more typically, at least about 80%; even more typically, at least about 85%; even more typically, at least about 90%; even more typically, at least about 95% receptor binding of the naturally occurring lipoproteins.

"Polynucleotide binding molecule" refers to those compounds that associate with polynucleotides, and the association is not sequence specific. For example, such molecules can (1) aid in neutralizing the electrical charge of polynucleotide, or (2) facilitate condensation of nucleotides, or (3) inhibit serum or nuclease degradation.

"Polycationic agent" refers generally to a polymer comprising positively-charged single units, although some non-positively charged units may be present in the polymer. The instant agents exhibit a net positive charge under physiologically relevant pH. Such agents are capable of neutralizing the charge of nucleic acids and can exhibit additional properties, such as condensation and/or serum protection of nucleic acids. Preferably, the agents comprises both amino acids and NSGs as monomeric units; also, preferred are agents comprising NSGs as monomeric units.

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"Physiologically relevant pH" varies somewhat between in vitro and in vivo applications. Typically, the physiological pH is at least 5.5; more typically, at least 6.0; even more typically, at least 6.5. Usually, physiologically relevant pH is no more than 8.5; more usually, no more than 8.0; even more usually, no more than 7.5.

"Polynucleotide" or "nucleic acid" refers to DNA, RNA, analogues thereof, peptide-nucleic acids, and DNA or RNA with non-phosphate containing nucleotides. Additionally, these nucleic acids may be single-stranded, double-stranded, or chimeric single- or double-stranded molecules.

The term "oligomer" includes polymers such as poly NSGs, produced by the submonomer process described herein and also in Zuckermann et al., supra. includes polymers, copolymers, and interpolymers of any length. More specifically, oligomers may comprise a single repeating monomer, two alternating monomer units, two or more monomer units randomly and/or deliberately spaced relative to each other. Regardless of the type of polyamide produced, the polyamide of the invention may be produced by the same general procedure which includes repeating a two-step or three step cycle wherein a new monomer unit is added in each cycle until an oligomer of desired length is obtained. The oligomer is preferably 2-100 monomers, more preferably 2-50, or 18-28 monomers or 24 to 48 monomers in length.

The term "frequency of uptake of polynucleotides into a cell" refers to an increase in the amount of polynucleotides actually taken up by a cell relative to the amount actually administered to the cell. The frequency of uptake of polynucleotides into a cell is increased if it is greater than the frequency of uptake of naked polynucleotides. For example, using *in vitro* transfection

methods, uptake of naked polynucleotides into mammalian cells is not usually detectable over background. Some frequency of uptake, however, can be detected when naked polynucleotides are delivered in vivo. The frequency of uptake in vivo and in vitro depends on the tissue type. The frequency of uptake can be measured by known methods for detecting the presence of polynucleotides, such as Northern, Southern, or Polymerase Chain Reaction (PCR) techniques.

Usually, a composition or compound is capable of increasing the frequency of polynucleotide uptake into a cell if it induces a frequency of uptake that is at least 10% greater than the frequency of naked polynucleotide uptake; more usually, at least 15% greater; even more usually, 20% greater; even more usually, at least 30%; and up to 40% to 100% greater, and even 1,000% and 10,000% greater.

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"Naked polynucleotides" refers to polynucleotides that are substantially free from any delivery vehicle that can act to facilitate entry into the cell. For example, polynucleotides are naked when free from any material which promotes transfection, such as liposomal formulations, charged lipids, such as Lipofectin® or precipitating agents such as $Ca_{3}(PO_{4})_{2}$.

"Effective amount to increase the frequency of polynucleotide uptake into cells" refers to an amount that induces a frequency of polynucleotide uptake into a cell that is at least 10% greater than the frequency of naked polynucleotide uptake; more usually, at least 15% greater; even more usually, 20% greater; even more usually, at least 30%; even more usually, at least 40%.

"Effective amount to neutralize nucleic acids" refers to the amount used to neutralize at least 10% of the electrical charge of the nucleic acid composition; more preferably; the amount refers to the amount used to neutralize at least 40%; even more preferably, the amount to neutralize 50% of the electrical charge; even more preferably, the amount to neutralize 60% of the electrical charge; even more preferably, the amount to neutralize 70% of the electrical charge; even more preferably, the amount to neutralize 80% of the electrical charge; and most preferably, at least 90% of the electrical charge of the nucleic acid composition of interest.

"Condensation of nucleic acids" occurs when the polycationic agent that is combined with nucleic acids, neutralizes the electrical charge of the nucleic acids and causes it to assume a reduced structure relative to uncomplexed nucleic acids. Preferably, condensation reduces the structure of nucleic acids to a size that can be internalized by structures present on cell surface membranes. Condensation can be measured by determining the charge of the nucleic acid/polycationic agent by gel electrophoresis, for example. Alternatively, an effective amount to condense nucleic acids can also be measured by the final size of the polycationic agent/nucleic acid complex.

"Effective amount to inhibit serum or nuclease degradation of nucleic acids" refers to the amount used to increase the half-life of the polynucleotide when exposed to serum and/or nucleases by at least 5 minutes as compared the uncomplexed nucleic acids; more preferably, the amount used to inhibit degradation by at least 10 minutes; even more preferably, the amount used to inhibit degradation by at least 30 minutes; even more preferably, the amount used to inhibit degradation by at least 45 minutes; even more preferably, the amount used to inhibit degradation by at least 60 minutes; even more preferably, the amount used to inhibit degradation by at least 90 minutes; and more preferably, the amount used to inhibit degradation by at least 120 minutes.

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A composition containing A is "substantially free of" B when at least 85% by weight of the total A+B in the composition is A. Preferably, A comprises at least about 90% by weight of the total of A+B in the composition, more preferably at least about 95% or even 99% by weight.

"Immunogenicity" refers to the ability of a given molecule or a determinant thereof to induce the generation of antibodies with binding capacity to the molecule upon administration in vivo, to induce a cytotoxic response, activate the complement system, induce allergic reactions, and the like. An immune response may be measured by assays that determine the level of specific antibodies in serum, by assays that determine the presence of a serum component that inactivates the polycationic agent/nucleic acid complex or conjugated gene delivery vehicle, or by other assays that measure a specific component or activity of the immune system. As discussed in more detail below, low immunogenicity may be established by these assays. The terms "low immunogenicity." "reduced

immunogenicity," "lowered immunogenicity" and similar terms are intended to be equivalent terms.

An "origin of replication" is a polynucleotide sequence that initiates and regulates replication of polynucleotides, such as an expression vector. The origin of replication behaves as an autonomous unit of polynucleotide replication within a cell, capable of replication under its own control. With certain origins of replication, an expression vector can be reproduced at a high copy number in the presence of the appropriate proteins within the cell. Examples of origins are the 2μ and autonomously replicating sequences, which are effective in yeast; and the viral T-antigen, effective in COS-7 cells.

General Methods and Detailed Description POLYNUCLEOTIDES

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Polynucleotides used in the instant invention can be used to express desired polypeptides, or can be, in themselves, therapeutic, such as ribozymes or antisense polynucleotides. Such polynucleotides can be used in *in vitro*, *ex vivo*, and *in vivo* applications.

Also, the polynucleotides of the invention can be vectors that express polypeptides, ribozymes, or antisense molecules. Vectors contain at least a promoter to initiate transcription operably linked to the coding region, ribozyme or antisense molecule. Other components that can be included in the vector are, for example: (1) a terminator sequence; (2) a sequence encoding a leader peptide to direct secretion; (3) a selectable marker; and (4) an origin of replication. An origin of replication is not necessary. The polynucleotides to be delivered can be either replicating or non-replicating. Other components can be added as desired and convenient.

The polynucleotides and methods of the invention can be utilized with any type of host cell. The choice of promoter, terminator, and other optional elements of an expression vector will depend on the host cell chosen. The invention is not dependent on the host cell selected. Convenience and the desired level of protein expression will dictate the optimal host cell. A variety of hosts for expression are known in the art and available from the American Type Culture Collection (ATCC) (Rockville, Maryland, U.S.A.). Suitable bacterial

hosts suitable include, without limitation: Campylobacter, Bacillus, Escherichia, Lactobacillus, Pseudomonas, Staphylococcus, and Streptococcus. Yeast hosts from the following genera may be utilized: Candida, Hansenula, Kluyveromyces, Pichia, Saccharomyces, Schizosaccharomyces, and Yarrowia. Aedes aegypti, Bombyx mori, Drosophila melanogaster, and Spodoptera frugiperda (PCT Patent Publication No. WO 89/046699; Carbonell et al., 1985, J. Virol. 56:153; Wright, 1986, Nature 321:718; Smith et al., 1983, Mol. Cell. Biol. 3:2156; and see generally, Fraser et al., 1989, In Vitro Cell. Dev. Biol. 25:225).

Useful mammalian cell types for *in vitro* applications include for example, those cell lines available from the American Type Culture Collection (ATCC), Chinese hamster ovary (CHO) cells, HeLa cells, baby hamster kidney (BHK) cells, monkey kidney cells (COS), human hepatocellular carcinoma cells (e.g., Hep G2), human embryonic kidney cells, baby hamster kidney cells, mouse sertoli cells, canine kidney cells, buffalo rat liver cells, human lung cells, human liver cells, mouse mammary tumor cells, as well as others.

Useful cell types for *in vivo* or *ex vivo* applications include, without limitation, any tissue type, such as muscle, skin, brain, lung, liter, spleen, blood, bone marrow, thymus, heart, lymph, bone, cartilage, pancreas, kidney, gall bladder, stomach, intestine, testis, ovary, uterus, rectum, nervous system, eye, gland, and connective tissue.

A. In vitro and Ex vivo Vectors

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The polynucleotides encoding the desired polypeptides or ribozymes, or antisense polynucleotides can be transcribed and/or translated using the following promoters and enhancers as examples. The examples include, without limitation: the 422(aP2) gene and the stearoyl-CoA desaturase 1 (SCD1) gene, which contains suitable adipocyte-specific promoters, as described in Christy et al., 1989, Genes Dev. 3:1323-1335. Synthetic non-natural promoters or hybrid promoters can also be used herein. For example, a T7T7/T7 promoter can be constructed and used, in accordance with Chen et al., 1994, Nucleic Acids Res. 22:2114-2120, where the T7 polymerase is under the regulatory control of its own promoter and drives the transcription of a polynucleotide sequence, which is

placed under the control of another T7 promoter. The primary determinant for the fat-specific expression is an enhancer located at about >5 kb upstream of the transcriptional start site, as described in Ross et al., 1990, Proc. Natl. Acad. Sci. USA.87.9590-9594 and Graves et al., 1991, Genes Dev. 5:428-437. Also suitable for use herein is the gene for the CCAAT/enhancer-binding protein C/EBPα, which is highly expressed when 3T3-L1 adioblast commit to the differentiation pathway and in mature post-mitotic adipocytes, as described in Birkenmeier et al., 1989, Gene Dev. 3:1146-1156. The recently isolated transcription factor PPARγ2, expressed exclusively in adipocyte tissues, as described in Tontonoz et al., 1994, Cell 79:1147-1156, can also be used herein.

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Typical promoters for mammalian cell expression include the SV40 early promoter, the CMV promoter, the mouse mammary tumor virus LTR promoter, the adenovirus major late promoter (Ad MLP), and the herpes simplex virus promoter, among others. Other non-viral promoters, such as a promoter derived from the murine metallothionein gene, will also find use in mammalian constructs. Expression may be either constitutive or regulated (inducible), depending on the promoter. Typically, transcription termination and polyadenylation sequences will also be present, located 3' to the translation stop codon. Preferably, a sequence for optimization of initiation of translation, located 5' to the coding sequence, is also present. Examples of transcription terminator/polyadenylation signals include those derived from SV40, as described in Sambrook et al., 1989, "Molecular Cloning, A Laboratory Manual," second edition, Cold Spring Harbor Press, Cold Spring Harbor, New York. Introns, containing splice donor and acceptor sites, may also be designed into the constructs of the present invention.

Enhancer elements can also be used herein to increase expression levels of the mammalian constructs. Examples include the SV40 early gene enhancer, as described in Dijkema et al., 1985, EMBO J. 4:761, and the enhancer/promoter derived from the long terminal repeat (LTR) of the Rous Sarcoma Virus, as described in Gorman et al., 1982b, Proc. Natl. Acad. Sci. USA 79:6777, and human cytomegalovirus, as described in Boshart et al., 1985, Cell 41:521. A leader sequence can also be present which includes a sequence encoding a signal peptide, to provide for the secretion of the foreign protein in mammalian cells. Preferably, there are processing sites encoded between the leader fragment and the gene of interest such that the leader sequence can be cleaved either in vivo or in vitro.

Other regulatory regions from viruses can be included in the polynucleotides of the instant invention to increase transcription and translation levels or increase the duration of transcription and translation. For example, the long terminal repeats of HIV can be included. Alternatively, the inverted terminal repeats of the Epstein Barr Virus can be used.

There exist expression vectors that provide for the transient expression in mammalian cells of DNA encoding the target polypeptide. In general, transient expression involves the use of an expression vector that is able to replicate efficiently in a host cell, such that the host cell accumulates many copies of the expression vector and, in turn, synthesizes high levels of a desired polypeptide encoded by the expression vector. Transient expression systems, comprising a suitable expression vector and a host cell, allow for the convenient positive identification of polypeptides encoded by cloned DNAs, as well as for the rapid screening of such polypeptides for desired biological or physiological properties. Thus, transient expression systems are particularly useful for purposes of identifying analogs and variants of the target polypeptide that have target polypeptide-like activity.

B. In vivo Vectors

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For delivery using viral vectors, any of a number of viral vectors can be used, as described in Jolly, 1994, Cancer Gene Therapy 1:1-64. For example, the coding sequence of a desired polypeptide or ribozymes or antisense molecules can be inserted into plasmids designed for transcription and/or translation in retroviral vectors, as described in Kimura et al., 1994, Human Gene Therapy 5:845-852, adenoviral vectors, as described in Connelly et al., 1995, Human Gene Therapy 6:185-193, adeno-associated viral vectors, as described in Kaplitt et al., 1994, Nature Genetics 6:148-153 and sindbis vectors. Promoters that are suitable for use with these vectors include the Moloney retroviral LTR, CMV promote and the mouse albumin promoter. Replication competent free virus can be produced and injected directly into the animal or humans or by transduction of an autologous cell ex vivo, followed by injection in vivo as described in Zatloukal et al., 1994, Proc. Natl. Acad. Sci. USA 91:5148-5152.

The polynucleotide encoding a desired polypeptide or ribozyme or antisense polynucleotide can also be inserted into plasmid for expression of the desired polypeptide in vivo. For in vivo therapy, the coding sequence can be delivered by direct injection into tissue, or via oral administration as an aerosol. Promoters suitable for use in this manner include endogenous and heterologous promoters such as CMV. Further, a synthetic T7T7/T7 promoter can be constructed in accordance with Chen et al., 1994, Nucleic Acids Res. 22:2114-2120, where the T7 polymerase is under the regulatory control of its own promoter and drives the transcription of polynucleotide sequence, which is also placed under the control of a T7 promoter. The polynucleotide can be injected in a formulation that can stablize the coding sequence and facilitate transduction thereof into cells and/or provide targeting, as described in Zhu et al., 1993, Science 261:209-211.

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Expression of the coding sequence of a desired polypeptide or replication of a ribozyme or antisense polynucleotide *in vivo* upon delivery for gene therapy purposes by either viral or non-viral vectors can be regulated for maximal efficacy and safety by use of regulated gene expression promoters as described in Gossen et al., 1992, Proc. Natl. Acad. Sci. USA 89:5547-5551. For example, the polynucleotide transcription and/or translation can be regulated by tetracycline responsive promoters. These promoters can be regulated in a positive or negative fashion by treatment with the regulator molecule.

For non-viral delivery of the coding sequence of the desired polypeptide, the sequence can be inserted into conventional vectors that contain conventional control sequences for high level expression.

C. Preferred Vector

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A preferred vector comprises: (1) an (EBV) origin of replication or a BKV (BK virus), a parvovirus, origin of replication; (2) a coding region for an EBV or BKV origin binding protein; (3) at least one inverted terminal repeat; (4) a promoter; (5) an enhancer; (6) a terminator; (7) optionally, a selectable marker.

Preferably, the orgin of replication is EBV ori p; more preferably, nucleotides 2623 to 4559 of SEQ ID NO:1 are utilized. The sequence is obtainable from vector pCEP4, commercially available from Invitrogen, San Diego, California, USA.

Preferably, the coding region encodes the EBV nuclear antigen A, which binds to EBV ori p; more preferably, the polynucleotide sequence is nucleotides 14 to 2594 of SEQ ID NO:1 are utilized. The sequence is obtainable from vector pCEP4, commercially available from Invitrogen, San Diego, California, USA.

Fragments and mutants of the preferred origin and binding protein capable of initiating replication of the vector in the desired host cell can be utilized. Preferably, the fragments and mutants will retain more than about 80% sequence identity with nucleotides 14 to 2594 or 2623 to 4559 of SEQ ID NO: 1 or fragment thereof; more typically, more than about 85%; even more typically, at least 90%. Preferably, these polynucleotides exhibit more than about 92% sequence identity with nucleotides 14 to 2594 or 2623 to 4559 of SEQ ID NO: 1 or fragment thereof; more preferably, more than about 94%; even more preferably, more than about 98%; even more preferably, more than about 98%; even more preferably, more than about 98%;

Preferably, the inverted terminal repeats are those sequences found in adenovirus (AV) or adeno-associated virus (AAV); more preferably, the inverted terminal repeats are those found in AAV; even more preferably, the polynucleotide sequence is 4938 to 5104 or 7189 to 7355 of SEQ ID NO: 1. The sequence of AAV is described in Samulski et al., 1987, J. Virol. 61:3096-3101.

Fragments and mutants of the preferred inverted terminal repeat capable of initiating replication of the vector in the desired host cell can be utilized. Preferably, the fragments and mutants will retain more than about 80% sequence identity with nucleotides 4938 to 5104 or 7189 to 7355 of SEQ ID NO: 1 or fragment thereof; more typically, more than about 85%; even more typically, at

least 90%. Preferably, these polynucleotides exhibit more than about 92% sequence identity with nucleotides 4938 to 5104 or 7189 to 7355 of SEQ ID NO: 1 or fragment thereof; more preferably, more than about 94%; even more preferably, more than about 96%; even more preferably, more than about 98%; even more preferably, more than about 99%.

Preferably, the cytomegalovirus enhancer/promoter is utilized; more preferably, the CMV promoter sequence is nucleotide sequence 5112 to 6734 of SEQ ID NO:1.

Mutants and fragments of the preferred enhancer and promoter capable of initiating transcription and/or translation can be utilized. Preferably, the fragments and mutants will retain more than about 80% sequence identity with nucleotides 5112 to 6734 of SEQ ID NO: 1 or fragment thereof; more typically, more than about 85%; even more typically, at least 90%. Preferably, these polynucleotides exhibit more than about 92% sequence identity with nucleotides 5112 to 6734 of SEQ ID NO: 1 or fragment thereof; more preferably, more than about 94%; even more preferably, more than about 96%; even more preferably, more than about 99%.

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A preferred terminator is the bovine growth hormone poly A sequence; more preferably, the polynucleotide sequence is nucleotide 6818 to 7050 of SEQ ID NO:1.

Mutants and fragments of the preferred terminator capable of terminating transcription and/or translation can be utilized. Preferably, the fragments and mutants will retain more than about 80% sequence identity with nucleotides 6818 to 7050 of SEQ ID NO: 1 or fragment thereof; more typically, more than about 85%; even more typically, at least 90%. Preferably, these polynucleotides exhibit more than about 92% sequence identity with nucleotides 6818 to 7050 of SEQ ID NO: 1 or fragment thereof; more preferably, more than about 94%; even more preferably, more than about 96%; even more preferably, more than about 98%; even more preferably, more than about 99%.

The sequence of the preferred vector is shown in SEQ ID NO:1. Polynucleotides encoding polypeptides, such as erythropoeitin or leptin, and ribozymes and antisense polynucleotides can be inserted into the vector.

D. Examples of Coding Regions, Ribozymes, and Antisense Molecules

The following are examples of coding regions, ribozymes, and antisense molecules that can be used to treat various indications in mammals. The nucleotide sequence of the genes of interest can be found, for example, in publically available databases, such as Genbank. Polynucleotides to be delivered can be used to treat viral infections or chronic pathogen infection.

1. Hemophilia

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Gene replacement by in vivo delivery of polynucleotides can be effective in treating hemophilia. The following are examples of polypeptides that can be encoded by the polynucleotides to be delivered: Factor VIII:C, mutants of Factor VIII:C, preferably those that are uncleavable. Also, useful to treat hemophilia are ribozyme and antisense polynucleotides as inhibitors of Tissue Factor Plasminogen Inhibitor (TFPI).

The routes of delivery for treating hemophilia include, for example, intravenous/intrahepatic injection, ex vivo transduction of stem cells or lymphocytes using retroviral vectors.

2. Treatment of Graft Versus Host Disease

In vivo delivery of polynucleotides encoding prodrugs can be used for direct ablation to treat graft versus host disease in, for example, leukemia bone marrow transplantation. Herpes thymidine kinase in conjunction with gancyclovir can be utilized for this purpose. Other examples of prodrugs are described in the cancer section.

The routes of delivery for treating graft versus host disease include, for example, ex vivo transduction of T-lymphocytes using retroviral vectors.

3. Vaccines

In vivo delivery of polynucleotides encoding a desired antigen can be utilized to induce an immune response. This response can include both cellular and humoral response. This type of vaccine can be used to treat cancer as well as infectious diseases. Further, such treatment can be either prophylactic or therpeutic immunotherapy.

Examples of infectious diseases include. Human Immunodeficiency Virus (HIV), Hepatitis A, B, C, etc., (HAV, HBV, HCV, etc.), Human Papiloma Virus (HPV), cytomegalovirus (CMV), herpes simplex 1 and 2 (HSV), etc. Preferred antigens include non-structural proteins 3, 4a, and 5b (NS3, NS4, and NS5b) of HCV; gB2 and gD2 of HSV; env and rev proteins of HIV.

Also, cancer antigens can be used in vaccines, for both therapeutic and prophylactic purposes.

The antigens can be presented in the context of Class I major histocompatibility antigens, or to induce a cellular cytotoxic T cell response, or to induce a humoral response comprising the synthesis of antibodies.

In addition, an antisense or ribozyme target to a immune suppressive molecule, IL-10, TGF- β , and CTLA-4, for example, can be useful to be administered with a vaccine.

The routes of delivery for vaccines include, for example, intramuscular injection, dendritic cell-based immunization, or oral immunization by both viral and non-viral vectors.

4. Diabetes Mellitus

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Diabetes is another indication that can be treated by *in vivo* delivery of a replacement gene. The following are examples of useful polypeptides to be encoded by the replacement gene: insulin, insulin-like growth factor I and II (IGF-I and II).

Also useful for treating diabetes are polynucleotides encoding IAS-L, found on the surface of B cells in the pancreas, to protect the cells from immune destruction.

The routes of delivery for treating diabetes include, for example, liver-directed, parotid-directed, pancreas-directed, salivary gland-directed using both viral and non-viral vectors.

5. Hyperlipidemia

Hyperlipidemia can be treated by *in vivo* delivery of the following

polynucleotides encoding apoproteins or lipoprotein receptors. A more extensive description of lipoproteins and apoproteins is provided below.

The routes of delivery for treating hyperlipidemia include, for example, liver-directed intravenous administration by both viral and non-viral vectors.

6. Myocardial Ischemia or Infarction

The following are examples of polynucleotides that are useful, when delivered *in vivo*, to treat myocardial ischemia or infarction:

polynucleotides encoding basic fibroblast growth factor (bFGF), fibroblast growth factor 5 (FGF-5) and IGF-I.

The routes of delivery for treating myocardial ischemia or infarction include, for example, intrapericardial delivery of viral vector or non-viral vectors.

7. Bowel Disease

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The following are examples of polynucleotides that can be delivered in vivo to treat bowel disease:

- (i) ribozymes or antisense polynucleotides as inhibitors of macrophage/inflammatory cell recruitment or activation, such as NFkB;
 - (ii) ribozymes or antisense polynucleotides to act as anti-apoptotic agents, such as inhibitors of interleukin 1b converting enzyme family;
 - (iii) polynucleotides encoding complement blockers, such as decay accelarting factor (DAF), membrane cofactor protein (MCP); and the fusions of DAF and MCP also known as CAB-2;
 - (iv) cyclooxygenase inhibitors;
 - (v) anti-proliferative agents, such as, ribozymes, antisense oligonucleotides, antibodies, protein, or peptides against c-myb, ras/raf, PI3 kinase, cyclins;
- 25 (vi) polynucleotides encoding suicide proteins/genes, such as, herpes thymidine kinase;
 - (vii) polynucleotides encoding replacement genes or proteins which maybe deficient or down regulated during the devleopment of inflammatory bowel disease.
- 30 (viii) polynucleotides encoding IkB.

8. Prostate Cancer and Benign Prostatic Hyperplasia

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The following polynucleotides can be delivered to treat prostate cancer and benign prostatic hyperplasia:

- (i) a polynucleotide encoding a pro-apoptotic agent, including for example, fas, fas ligand, fadd, fap-1, tradd, faf, rip, reaper, apoptin, interleukin-2 converting enzyme;
 - (ii) a polynucleotide encoding an anti-angiogenic agent, including, for example, bFGF soluble receptor and fragments, angiostatin, transforming growth factor- β (TGF- β), interferon- α (IFN α), proliferin-related protein, a urokinase plasminogen activator receptor antagonist, platelet factor 4 (PF4), thrombospondin, a tissue inhibitor of metalloproteinase, and prolactin;
 - (iii) a polynucleotide encoding a immunomodulating agent including, for example, interleukin-2 (IL-2), IFN α , IFN β , IFN γ , granulocyte macrophage-colony stimulating factor (GM-CSF), and macrophage-colony stimulating factor (M-CSF);
- (iv) a ribozyme or antisense polynucleotide as an antiproliferative agent including, for example, an inhibitor of a signal transduction pathway, for example, an inhibitor of a signal transduction pathway mediated by myb, ras, ras superfamily, raf, phosphoinositol (PI3-kinase), a phosphotyrosine binding (PTB) domain, a SRC homology-2 (SH2) domain, a SRC homology-3 (SH3) domain, a plextrin homology (PH) domain, JUN kinase, and a stress activated kinase, signaling inositol phosphatases; and an inhibitor of a cyclin;
- (v) a ribozyme or antisense polynucleotide as an inhibitor of a growth factor or inhibitor of a receptor of a growth factor, including, for example, epidermal growth factor (EGF), TGF-α, FGF, TGF-β, platelet derived growth factor (PDGF), keratinocyte growth factor (KGF), or any prostate cell specific growth factor;
- (vi) a polynucleotide encoding a tumor suppressor gene or a gene downregulated during the onset of a hyperplastic condition in the prostate; and
- (vii) an antisense or ribozyme target to a immune suppressive molecule, IL-10, TGF- β , and CTLA-4, for example.

9. Anemia, Leukopenia, and Thrombocytopenia

Anemia can be treated by *in vivo* delivery of a polynucleotide encoding erythropoietin, GM-CSF-, G-CSF, M-CSF, and thrmobopoietin, for example. Examples of delivery routes for this indication include without limitation: livertargeted intravenous administration of viral vectors and non-viral vectors. See the Examples below.

10. Cardiomyopathy

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The following are examples of polynucleotides that can be delivered in vivo to treat cardiomyopathy: polynucleotides encoding, IGF-1, L-amino acid decarboxylas, inhibitors of β adrenergic receptor kinases (BARK), troponin, and β adrenergic receptors.

Examples of delivery routes for this indication include, without limitation, pericardial expression of IGF-1, and for the other genes, intramycardial injection or myocardial trageting via intracoronary injection or intrapericardial administration of viral vectors or non-viral vectors.

11. Rheumatoid Arthritis

The following are examples of polynucleotides that can be delivered in vivo to treat rheumatoid arthritis, polynucleotides encoding a prodrug, such as herpes thymidine kinase, MMP inhibitors, fas, and pro-apoptotic proteins, described above, and interleukin-1 receptor A, interleukin-10, IkB.

Also, antisense and ribozyme polynucleotides as inhibitors of NFkB.

Examples of delivery routes for this indication include, without limitation, intraarticular injection of viral and non-viral vectors.

12. Osteoarthritis and Psoriasis

The following are examples of polynucleotides that can be delivered in vivo to treat osteoarthritis and psoriasis: polynucleotides encoding IGF-1; ribozyme and antisense polynucleotides as inhibitors of metalloproteinase inhibitors.

Also, the following are examples of polynucleotides that can be delivered in vivo to treat osteoarthritis and psoriasis, polynucleotides encoding a prodrug,

such as herpes thymidine kinase, MMP inhibitors, fas, and pro-apoptotic proteins, described above, and interleukin-1 receptor A, interleukin-10, IkB.

Also, antisense and ribozyme polynucleotides as inhibitors of NFkB.

Examples of delivery routes for this indication include, without limitation, intraarticular injection.

13. Restenosis

The following are examples of polynucleotides that can be delivered in vivo to treat restenosis:

- (i) polynucleotides encoding a prodrug, such as thymidine kinase,
 other examples are described in the cancer section;
 - (ii) polynucleotides encoding tissue factor plasminogen inhibitor (TFPI);
 - (iii) polynucleotides encoding c-myb rbz, c-ras rbz,
 - (iv) polynucleotides encoding pro-apoptotic agents, described above;
- (v) polynucleotides encoding IκB.

Examples of delivery routes for this indication include, without limitation, intracoronary delivery of viral and non-viral vectors.

14. Cancer

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The gene delivery vectors of the invention are useful in delivering
therapeutic genes for treatment of hyperproliferative disorders, including
malignancy, for treatment of infectious disease and for treatment of
inflammatory diseases, including autoimmune disease. For instance, the gene
therapy vectors can be used to express cytokines or proteins that convert an
inactive or partially active prodrug into an active drug. In many cases,
conversion of the prodrug into its active form results in a compound with
cytolytic activity.

a. Prodrug converting enzymes

A number of "suicide genes" which encode different proteins useful in prodrug conversion can be used in the instant invention. For instance, nucleoside kinases such as thymidine kinase are particularly useful. In particular, the HSV-

TK system has important advantages for anti-tumor cell therapy. See PCT publication number WO 91/02805 entitled "Recombinant Retroviruses Delivering Vector Constructs to Target Cells" and PCT publication number WO 95/14014091 entitled "Compositions and Methods for Utilizing Conditionally

Lethal Genes" for a description of treatment of cancer and other diseases by gene 5 delivery vectors expressing thymidine kinase and other prodrug converting enzymes. HSV-TK transduced tumor cells can mediate a significant bystander killing effect on untransduced neighboring cells in vitro and in vivo (Moolten et al., supra., Freeman et al., 1993, Cancer Res. 53:5274), most commonly as a result of transfer to the toxic ganciclovir metabolite, GCV triphosphate, between adjacent cells through intercellular gap junctions (Bi et al., 1993, Human Gene Therap. 4:725). Endothelial cells in capillary walls are connected by gap

junctions, so a dramatic bystander effect created by GCV-triphosphate transfer between neighboring endothelial cells and the massive amplification effects of the clotting cascade and the tumor to endothelial cell ratio could ensue 15 (Denekamp et al., 1986, Cancer Topics 6:6; Denekamp et al., 1984, Prog. Appl. Microcir. 4:28). Recent evidence suggests that the occasional transduction of tumor endothelial cells during intralesional therapy with HSV-TK retroviral vectors may account for a significant component of the antitumor activity of the

vectors (Ram et al., 1994, J. Neurosurg. 81:256). In addition, the suicide gene is only conditionally cytotoxic to the target cells (i.e. only when GCV is given). Consequently, an ex vivo administration method can be be utilized. For example, in this type of protocol, endothelial cells may be isolated from tumor biopsies (Medzelewski et al., 1994, Cancer Res. 54:336), induced to proliferate with appropriate mitogens (Ferrara et al., supra.) and transduced with TK in vitro.

Transplanted EC become incorporated into the neovasculature in days to weeks after intratumoral injection (Lal et al., 1994, Cancer Gene Therap.1:322), so GCV treatment would follow a suitable 'lag phase' to allow the transduced EC to integrate functionally in to the tumor vasculature. The two-step enzyme-prodrug system offers greater flexibility of delicate clinical management, because cessation of GCV infusion in the event of (potentially very serious) complications arising from damage to normal EC, would block toxicity without

the need to block transgene activity in situ.

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A number of alternative 'suicide genes' in addition to thymidine kinase may also be useful for cancer gene therapy (Moolten et al., supra.). Introduction of the bacterial cytosine deaminase gene (Huber et al., 1993, Cancer Res. 53:4619) into tumor cells confers sensitivity to the antifungal agent 5-

- fluorocytosine (5-FC). Cytosine deaminase converts 5-FC to 5-fluorouracil (5-FU, Nishiyama et al., 1985, Cancer Res. 45:1753). Since 5-FU is commonly used chemotherapeutic drug for breast cancer, several groups have developed cytosine deaminase-based 'suicide gene' therapy models for this disease. Tumor specificity may be further increased by introducing the c-erbB2
- promoter/enhancer elements 5' to the cytosine deaminase gene, so that the therapeutic transgene is preferentially transcribed in c-erbB2-overexpressing breast tumor cells (Harris et al., 1994, Gene Therap. 1:170). Alkaline phosphatase has been widely explored as prodrug-activating enzyme in the related field of antibody directed enzymne-prodrug therapy (ADEPT). This enzyme has the advantage that it can activate a wide range of phosphorylated
- derivatives of anticancer agents (e.g. mitomycin C, etoposide, etc.) that cannot cross cell membranes until the charged phosphate group is cleaved off, so a single enzyme could generate de novo a cocktail of chemotherapeutic agents within the tumor mass (Senter et al., 1993, Bioconjugate Chem. 4:3). Other suicide genes may encode a polypeptide or polypeptides (with a corresponding non-cytotoxic agent) such as Herpes Simpex virus thymidine kinase (gancyclovir or acyclovir), Varicella Zoster virus thymidine kinase (6 methoxypurine arabino

nucleoside; Huber et al., 1991, Proc. Natl. Acad. Sci. USA 88:8039), E. coli

- cytosine deaminase (fluorouracil; Mullen et al., 1992, Proc. Natl. Acad. Sci.

 25 USA. 89:33), E. coli xanthine-guanine phophoribosyl transferase (thioxanthine; Beshard et al., 1987, Mol. Cell Biol. 7:4139), E. coli or Leishmania purine nucleotide phosphorylase (various nontoxic purine deoxyadenosine, adenosine, deoxyguanosine, or guanosine derivatives (Koszalka and Krenitsky, 1979, J. Biol Chem 254:8185, 1979; Sorscher et al., 1994, Gene Therapy 1:233), cytochrome pla50 2B1 or cytochrome pd50 reductors (social contents of the cytochrome pd50 reductors (social contents).
 - pla50 2B1 or cytochrome p450 reductase (e.g., 3amino-1,2,4 benzotriazine 1,4-dioxide (Walton et al.,1992, Biochem. Pharmacol. 44:251), cell surface alkaline phosphatase (e.g., etoposide monophosphate; Senter et al., 1988, Proc. Natl. Acad. Sci. USA 85:4842, 1988), nitroreductase (e.g., metronidazole or

nitrofurantoin: Hof et al., 1988, Immunitat und Infektion 16:220), N-deoxyribosy transferase (1-deazapurine; Betbeder et al., 1989; Nucleic Acids Res 17:4217), pyruvate ferrodoxin oxidoreductase (metronidazol; Upcroft et al., 1990, Int. J. Parasitolog, 20:489), carboxypepidase G2 (aminoacylate nitrogen mustards;

5 Antoniw et al., 1990, Brit J. Cancer 62:909), carboxypeptidase A (methotrexate alpha alanine; Haenseler et al., 1992, Biochemistry 31:891), • lactamase (cephalosporin derivatives; Meyer et al., 1993, Cancer Res. 53:3956; and Vradhula et al., 1993, Bioconjugate Chemistry 4:334), Actinomycin D synthetase complex (synthetic pentapeptide lactone precursors; Katz et al., 1990, J. Antibiotics 43:231), and •-glucuronidase (various glucuronide derivatives of toxic drugs such as doxorubicin; Bosslet et al., 1994, Cancer Res. 54:2151; Haeberlin et al., 1993, Pharmaceutical Res. 10:1553).

Any of a variety of other enzymes which convert inactive prodrugs into active drugs and known to those of skill in the art can also used in the gene delivery vehicles of the invention. For example, see PCT publication number WO 95/14014091 entitled "Compositions and Methods for Utilizing Conditionally Lethal Genes", and European Patent publication number EP90309430, entitled "Molecular Chimeras Useful for Cancer Therapy—Comprising Regulatory Sequences and heterologous enzyme, e.g. Varicella Zoster Virus Thymidine Kinase" for a description of additional prodrug/enzyme systems useful for gene therapy. As an additional example, see PCT Patent Publication No. WO 95/13095 entitled "New Prodrugs and Enzyme Targeting Molecule Conjugates—Useful in Antibody Direct Enzyme Prodrug Therapy of e.g. Viral Infections".

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A variety of tumors may be targeted for treatment by the gene delivery vehicles of the invention. In general, solid tumors are preferred, although leukemias and lymphomas may also be treated if they have developed a solid mass, or if suitable tumor associated markers exist such that the tumor cells can be physically separated from nonpathogenic normal cells. Representative examples of suitable tumors include melanomas, colorectal carcinomas, lung carcinomas (including large cell, small cell, squamous and adeno-carcinomas), renal cell carcinomas and breast adeno-carcinomas. Gene delivery vehicles expressing thymidine kinase and other prodrug converting enzymes are also

useful in the treatment of autoimmune diseases including rheumatoid arthritis, osteoarthritis and graft vs. host disease. See e.g. PCT Patent Publication No. WO 95/14091, entitled "Compositions and Methods for Utilizing Conditionally Lethal Genes," for a description of treatment of disease with gene therapy vectors expressing prodrug converting enzymes.

b. Cytokines

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A variety of polynucleotides encoding cytokines and immune system modulators can be delivered by the gene delivery vehicles of the invention for treatment of a number of different disorders. Representative examples include cytokines, such as IL-1, IL-2 (Karupiah et al., 1990, J. Immunology 144:290-298; Weber et al., 1987, J. Exp. Med. 166:1716-1733; Gansbacher et al., 1990, J. Exp. Med. 172:1217-1224; U.S. Patent No. 4,738,927), IL-3, IL-4 (Tepper et al., 1989, Cell 57:503-512; Golumbek et al., 1991, Science 254:713-716, 1991; U.S. Patent No. 5,017,691), IL-5, IL-6 (Brakenhof et al., 1987, J. Immunol. 139:4116-4121; WO 90/06370), IL-7 (U.S. Patent No. 4,965,195), IL-8, IL-9, IL-10, IL-11, IL-12, IL-13 (Cytokine Bulletin, Summer 1994), IL-14 and IL-15, particularly IL-2, IL-4, IL-6, IL-12, and IL-13, alpha interferon (Finter et al., 1991, Drugs 42(5):749-765; U.S. Patent No. 4,892,743; U.S. Patent No. 4,966,843; WO 85/02862; Nagata et al., 1980, Nature 284:316-320; Familletti et al., 1981, Methods in Enz. 78:387-394; Twu et al., 1989, Proc. Natl. Acad. Sci. USA 86:2046-2050; Faktor et al., 1990, Oncogene 5:867-872), beta interferon (Seif et al., 1991, J. Virol. 65:664-671), gamma interferons (Radford et al., The American Society of Hepatology 2008-2015, 1991; Watanabe et al., PNAS 86:9456-9460, 1989; Gansbacher et al., 1990, Cancer Research 50:7820-7825; Maio et al., 1989, Can. Immunol. Immunother. 30:34-42; U.S. Patent No. 4,762,791; U.S. Patent No. 4,727,138), G-CSF (U.S. Patent Nos. 4,999,291 and 4,810,643), GM-CSF (WO 85/04188), tumor necrosis factors (TNFs) (Jayaraman et al., 1990, J. Immunology 144:942-951), CD3 (Krissanen et al., 1987, Immunogenetics 26:258-266, 1987), ICAM-1 (Altman et al., 1989, Nature 338:512-514; Simmons et al., 1988, Nature 331:624-627), ICAM-2, LFA-1, LFA-3 (Wallner et al., 1987, J. Exp. Med. 166(4):923-932), MHC class I molecules, MHC class II molecules, B7.1-.3, 2-microglobulin (Parnes et al.,

1981, Proc. Natl. Acad. Sci. 78:2253-2257), chaperones such as calnexin, MHC linked transporter proteins or analogs thereof (Powis et al., 1991, Nature 354:528-531, 1991).

Genes encoding any of the cytokine and immunomodulatory proteins described herein can be expressed in a gene delivery vehicle of the invention. Other forms of these cytokines which are known to those of skill in the art can also be used. For instance, nucleic acid sequences encoding native IL-2 and gamma-interferon can be obtained as described in U.S. Patent Nos. 4,738,927 and 5,326,859, respectively, while useful muteins of these proteins can be obtained as described in U.S. Patent No.4,853,332. As an additional example, nucleic acid sequences encoding the short and long forms of mCSF can be obtained as described in US Patent Nos. 4, 847,201 and 4,879,227, respectively.

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encoding Interleukin-6).

Other nucleic acid molecules that encode cytokines, as well as other nucleic acid molecules that are advantageous for use within the present invention, may be readily obtained from a variety of sources, including, for 15 example, depositories such as the American Type Culture Collection (ATCC, Rockville, Maryland), or from commercial sources such as British Bio-Technology Limited (Cowley, Oxford England). Representative examples include BBG 12 (containing the GM-CSF gene coding for the mature protein of 127 amino acids), BBG 6 (which contains sequences encoding gamma 20 interferon), ATCC No. 39656 (which contains sequences encoding TNF), ATCC No. 20663 (which contains sequences encoding alpha interferon), ATCC Nos. 31902, 31902 and 39517 (which contains sequences encoding beta interferon), ATCC No 67024 (which contains a sequence which encodes Interleukin-1b), ATCC Nos. 39405, 39452, 39516, 39626 and 39673 (which contains sequences 25 encoding Interleukin-2), ATCC Nos. 59399, 59398, and 67326 (which contain sequences encoding Interleukin-3), ATCC No. 57592 (which contains sequences encoding Interleukin-4), ATCC Nos. 59394 and 59395 (which contain sequences

Gene delivery vehicles expressing the above cytokines are useful in the treatment of a variety of disorders. For example, see PCT publication number

encoding Interleukin-5), and ATCC No. 67153 (which contains sequences

US94/02951 entitled "Compositions and Methods for Cancer Immunotherapy" for a description of gene therapy treatment of malignancy.

15. Neurological Disorders and Diseases

Polynucleotides encoding tyrosine hydroxylase can be useful in treating

Parkinson disease.

For stroke or any acute brain injuries, polynucleotides encoding IGF-1, bFGF, vascular endothelial growth factor (VEGF) are useful.

16. Pulmonary Disorders

For treating emphysema, polynucleotides encoding $\alpha 1$ -anti-trypsin are useful.

For treating lung fibrosis, polynucleotides encoding superoxide dismutase (SOD) are useful.

For treating cystic fibrosis, polynucleotides encoding CFTR are useful.

15 ADDITIONAL AGENTS

Additional agents can be included with the desired polynucleotides to be delivered. These additional agents can facilitate endocytosis of the desired nucleic acids or aid binding of the nucleic acids to the cell surface or both, for example.

20 A. Polypeptides

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One example are polypeptides which include, without limitation: asialoorosomucoid (ASOR); transferrin; asialoglycoproteins; antibodies; antibody fragments; ferritin; interleukins; interferons, granulocyte, macrophage colony stimulating factor (GM-CSF), granulocyte colony stimulating factor (G-CSF), macrophage colony stimulating factor (M-CSF), stem cell factor and erythropoietin. Viral antigens, such as envelope proteins, can also be used. Also, proteins from other invasive organisms, such as the 17 amino acid peptide from the circumsporozoite protein of plasmodium falciparum known as RII.

B. Hormones, Vitamins, Etc.

Other groups that can be included are, for example: hormones, steroids, androgens, estrogens, thyroid hormone, or vitamins, folic acid.

C. Polyalkylenes, Polysaccharides, Etc.

Polyalkylene glycols can be included with the desired polynucleotides. In a preferred embodiment, the polyalkylene glycol is polyethlylene glycol. In addition, mono-, di-, or polysaccarides can be included. In a preferred embodiment of this aspect, the polysaccharide is dextran or DEAE-dextran. Also, chitosan and poly(lactide-co-glycolide)

D. Lipids and Liposomes

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The desired polynucleotide can also be encapsulated in lipids or packaged in liposomes prior to delivery to the subject or to cells derived therefrom.

Lipid encapsulation is generally accomplished using liposomes which are able to stably bind or entrap and retain nucleic acid. The ratio of condensed polynucleotide to lipid preparation can vary but will generally be around 1:1 (mg DNA:micromoles lipid), or more of lipid. For a review of the use of liposomes as carriers for delivery of nucleic acids, see, Hug and Sleight, 1991, <u>Biochim</u>. <u>Biophys</u>. <u>Acta</u>. <u>1097</u>:1-17; Straubinger *et al.*, in METHODS OF ENZYMOLOGY (1983), Vol. 101, pp. 512-527.

Liposomal preparations for use in the instant invention include cationic (positively charged), anionic (negatively charged) and neutral preparations.

Cationic liposomes have been shown to mediate intracellular delivery of plasmid DNA (Felgner et al., 1987, Proc. Natl. Acad. Sci. USA 84:7413-7416); mRNA (Malone et al., 1989, Proc. Natl. Acad. Sci. USA 86:6077-6081); and purified transcription factors (Debs et al., 1990, J. Biol. Chem. 265:10189-10192), in functional form.

Cationic liposomes are readily available. For example, N[1-2,3-dioleyloxy)propyl]-N,N,N-triethylammonium (DOTMA) liposomes are available under the product line Lipofectin⁶, from GIBCO BRL, Grand Island, NY. (See, also, Felgner et al., 1987, Proc. Natl. Acad. Sci. USA 84:7413-7416). Other commercially available liposomes include transfectace (DDAB/DOPE) and

DOTAP/DOPE (Boerhinger). Other cationic liposomes can be prepared from readily available materials using techniques well known in the art. See, e.g., Szoka et al., 1978, Proc. Natl. Acad. Sci. USA 75:4194-4198; PCT Publication No. WO 90/11092 for a description of the synthesis of DOTAP (1,2-bis(oleoyloxy)-3-(trimethylammonio)propane) liposomes.

Similarly, anionic and neutral liposomes are readily available, such as from Avanti Polar Lipids (Birmingham, AL), or can be easily prepared using readily available materials. Such materials include phosphatidyl choline, cholesterol, phosphatidyl ethanolamine, dioleoylphosphatidyl choline (DOPC), dioleoylphosphatidyl glycerol (DOPG), dioleoylphoshatidyl ethanolamine (DOPE), among others. These materials can also be mixed with the DOTMA and DOTAP starting materials in appropriate ratios. Methods for making liposomes using these materials are well known in the art.

unilamellar vesicles (SUVs), or large unilamellar vesicles (LUVs). The various liposome-nucleic acid complexes are prepared using methods known in the art. See, e.g., Straubinger et al., in METHODS OF IMMUNOLOGY (1983), Vol. 101, pp. 512-527; Szoka et al., 1978, Proc. Natl. Acad. Sci. USA 75:4194-4198; Papahadjopoulos et al., 1975, Biochim. Biophys. Acta 394:483; Wilson et al., 1979, Cell 17:77; Deamer and Bangham, 1976, Biochim. Biophys. Acta 443:629; Ostro et al., 1977, Biochem. Biophys. Res. Commun. 76:836; Fraley et al., 1979, Proc. Natl. Acad. Sci. USA 76:145); Fraley et al., 1980, J. Biol. Chem. 255:10431; Szoka and Papahadjopoulos, 1978, Proc. Natl. Acad. Sci. USA 75:145; and Schaefer-Ridder et al., 1982, Science 215:166.

E. Lipoproteins

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In addition, lipoproteins can be included with the polynucleotide to be delivered. Examples of lipoproteins to be utilized include: chylomicrons, HDL, IDL, LDL, and VLDL. Mutants, fragments, or fusions of these proteins can also be used. Also, modifications of naturally occurring lipoproteins can be used, such as acetylated LDL. These lipoproteins can target the delivery of polynucleotides to cells expressing lipoprotein receptors. Preferably, if

lipoproteins are including with the polynucleotide to be delivered, no other targeting ligand is included in the composition.

If lipoproteins are included with the desired polynucleotides to be delivered, preferably, the composition comprises: (1) lipoprotein; (2) polynucleotide; and (3) a polynucleotide binding molecule.

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Naturally occurring lipoproteins comprise a lipid and a protein portion. The protein portion are known as apoproteins. At the present, apoproteins A, B, C, D, and E have been isolated and identified. At least two of these contain several proteins, designated by Roman numerals, AI, AII, AIV; CI, CII, CIII.

A lipoprotein can comprise more than one apoprotein. For example, naturally occurring chylomicrons comprise A, B, C, and E, over time these lipoproteins lose A and acquire C and E apoproteins. VLDL comprises A, B, C, and E apoproteins, LDL comprises apoprotein B; and HDL comprises apoproteins A, C, and E.

The amino acids of these apoproteins are known and are described in, for example, Breslow, 1985, Annu Rev. Biochem 54:699; Law et al., 1986, Adv. Exp Med. Biol. 151:162; Chen et al., 1986, J Biol Chem 261: 12918; Kane et al., 1980, Proc Natl Acad Sci USA 77:2465; and Utermann et al., 1984, Hum Genet 65:232.

Lipoproteins contain a variety of lipids including, triglycerides, cholesterol (free and esters), and phopholipids. The composition of the lipids varies in naturally occurring lipoproteins. For example, chylomicrons comprise mainly triglycerides. A more detailed description of the lipid content of naturally occurring lipoproteins can be found, for example, in Meth. Enzym. 128 (1986). The composition of the lipids are chosen to aid in conformation of the apoprotein for receptor binding activity. The composition of lipids can also be chosen to facilitate hydrophobic interaction and association with the polynucleotide binding molecule.

Naturally occurring lipoproteins can be isolated from serum by ultracentrifugation, for instance. Such methods are described in <u>Meth. Enzy.</u>, supra; Pitas et al., 1980, <u>J. Biochem. 255</u>:5454-5460; and Mahey et al., 1979, <u>J. Clin. Invest 64</u>:743-750.

Lipoproteins can also be produced by *in vitro* or recombinant methods by expression of the apoprotein genes in a desired host cell. See, for example, Atkinson et al., 1986, Annu Rey Biophys Chem 15:403, and Radding et al., 1958, Biochim. Biophys Acta 30:443.

Lipoproteins can also be purchased from commercial suppliers, such as Biomedical Techniologies, Inc., Stoughton, Massachusetts, USA.

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Mutants, fragments and fusion of the naturally occurring apoproteins are useful for delivery of polynucleotides. These polypeptides will retain more than about 80% amino acid identity; more typically, more than about 85%; even more typically, at least 90%. Preferably, these polypeptides will exhibit more than about 92% amino acid sequence identity with naturally occurring lipoproteins or fragment thereof; more preferably, more than about 94%; even more preferably, more than about 96%; even more preferably, more than about 98%; even more preferably, more than about 98%; even more

Such mutants, fragments and fusions can be constructed by altering the polynucleotides encoding the desired lipoproteins by recombinant DNA techniques. See, for example, Sambrook et al., (1989) Molecular Cloning, A Laboratory Manual, 2d edition (Cold Spring Harbor Press, Cold Spring Harbor, New York). These polynucleotides can be inserted into expression vectors and host cells can be utilized to produce the desired apoprotein.

In addition, naturally occurring lipoproteins, mutants, fragments, and fusions can be chemically altered. For example, acetylated LDL has biological activity. See, for example, Nagelkerke et al., 1983, J. Biol. Chem. 258(20):12221-12227; Weisgraber et al., 1978, J. Biol. Chem. 253:9053-9062;

Voyta et al., 1984, J. Cell Biol. 99:2034-2040; Goldstein et al., 1979, Proc. Natl. Acad. Sci. USA 76:333-337; and Pitas, 1981, Arterosclerosis 1:177-185.

Chemically modified lipoproteins can also be purchased from commercial suppliers, such as Biomedical Techniologies, Inc., Stoughton, Massachusetts, USA.

All of these polypeptides exhibit receptor binding properties of naturally occurring lipoproteins. Usually, such polypeptides exhibit at least about 20% receptor binding of naturally occurring lipoproteins. More typically, the polypeptides exhibit at least about 40%, even more typically the polypeptides

exhibit at least about 60%; even more typically, at least about 70%; even more typically, at least about 80%; even more typically, at least about 85%; even more typically, at least about 95% receptor binding of the naturally occurring lipoproteins.

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Typically, lipoproteins are present in an amount effective to increase the frequency of incorporation of polynucleotides into a cell. Such an amount is sufficient to increase the frequency of incorporation of polynucleotides into a cell by at least 10%, compared to the frequency of incoporation of naked polynucleotides; more usually, at least 15%; even more usually, 20%; even more usually, at least 30%. The increase can be between 40 to 100%, and even 1000% and 10000% increase.

"Polynucleotide binding molecule" refers to those compounds that associate with polynucleotides, and the association is not sequence specific. For example, such molecules can (1) aid in neutralizing the electrical charge of polynucleotide, or (2) facilitate condensation of nucleotides, or (3) inhibit serum or nuclease degradation. Optionally, polynucleotide binding molecules can interact with lipoproteins by either hydrophobic association or by charge. Polynucleotide binding molecules include, without limitation, polypeptides, mineral compounds, vitamins, etc.

Examples of polynucleotide binding molecules include: polylysine, polyarginine, polyornithine, and protamine. Examples of organic polycations include: spermine, spermidine, and purtrescine. Other examples include histones, protamines, human serum albumin, DNA binding proteins, non-histone chromosomal proteins, coat proteins from DNA viruses, such as \$\phi X174\$, transcriptional factors also contain domains that bind DNA and therefore may be useful as nucleic aid condensing agents. Briefly, transcriptional factors such as C/CEBP, c-jun, c-fos, AP-1, AP-2, AP-3, CPF, Prot-1, Sp-1, Oct-1, Oct-2, CREP, and TFIID contain basic domains that bind DNA sequences.

Examples of other positively charged moieties include polybrene, DEAE-dextran, and cationic lipids. Useful cationic lipids and liposomes are described above. Lipids and liposomes are not used in this aspect of the invention to encapsulate both polynucleotide and lipoprotein. The lipoprotein must be exposed to bind the its cell surface receptor.

Other synthetic compounds that are capable of binding negatively charged polynucleotides are useful, such as polymers of N-substituted glycines and others, as described below.

In a composition with a lipoprotein, the polynucleotide binding molecule can be present in an amount effective to neutralize the polynucleotide. However, the polynucleotide binding molecule also can be in excess of an effective amount to neutralize the polynucleotide to be delivered. Such an excess can produce a net positive electrical charge when complexed with the polynucleotides to be delivered. The positively charged complex can then interact with lipoproteins that comprise negatively charged lipids, such as phospholipids.

Typically, the polynucleotide binding molecule is in excess when the amount is 10% greater than the amount to neutralize the polynucleotide charge; more typically, the amount is 50% greater; even more typically, 100% greater; even more typically, 200% greater; even more typically, 200% greater; even more typically, 20,000% greater; even more typically, 20,000% greater; even more typically, 25,000% greater; even more typically, 25,000% greater; even more typically, 30,000% greater; even more typically, more than 40,000% greater than the amount effective to neutralize the electrical charge of the desired polynucleotide.

20 POLYCATIONIC AGENTS

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Polycationic agents can be included, with or without lipoprotein, in a composition with the desired polynucleotide to be delivered.

Functional properties

A. Net Positive Charge

25 Polycationic agents typically exhibit a net positive charge at physiological relevant pH and are capable of neutralizing the electrical charge of nucleic acids to facilitate delivery to a desired location. These agents have both in vitro, ex vivo, and in vivo applications. For example, these polycationic agents can be used to transfect cells used to produce recombinant proteins.

Alternatively, the instant polycationic agents can be used to deliver nucleic acids to a living subject either intramuscularly, subcutaneously, etc.

Physiological relevant pH varies somewhat between *in vitro* and *in vivo* applications. Typically, physiological pH is at least 5.5; more typically, at least 6.0; even more typically, at least 6.5. Usually, physiologically relevant pH is no more than 8.5; more usually, no more than 8.0; even more usually, no more than 7.5.

Preferably, the isoelectric point of the instant polycationic agents to neutralize nucleic acids is at least 9.

B. Non-Toxicity and Non-Immunogenic Properties

The composition of the polycationic agents of the invention will exhibit the toxicity and immunogenic properties desired. *In vitro* cell culture will have different immunogenic constraints than *in vivo* mammalian applications.

The instant polycationic agents can be easily tested for toxicity. For example, the agents can be added to medium for cells used in the *in vitro* assays, such as cos-7, Chinese Hamster Ovary cells, etc. Alternatively, the agents can be tested in standard animal tests for safety.

C. Condensation Properties

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Due to the electric charge, a subset of these polycationic agents are capable of condensing the desired nucleic acids to a compact size to facilitate delivery. Typically, condensation "collapses" polynucleotides or nucleic acids into macromolecular structures, commonly into a toroid form. The smaller size of condensed nucleic acids eases delivery by facilitating, for example, packaging nucleic acids into liposomes and/or reducing exposure to proteases and/or nucleases.

The condensed nucleic acids exhibit different properties compared to "relaxed" nucleic acids, such as (1) a decrease in intercalation of ethidium bromide or other intercalating dye or (2) a reduced mobility in gel electrophoresis. Thus, condensation can be measured by at least two different assays, an intercalating dye assay or a band shift assay.

One type of intercalating dye assay uses ethidium bromide. In this assay, test nucleic acids, conveniently plasmid DNA, are mixed with polycationic agent in a ratio from about 1:1 to a 1:50 weight/weight ration of plasmid to condensing

agent. Following incubation, ethidium bromide is added to the reaction to a final concentration of 1 µg/mL. If a nucleic acid such as RNA is used as the test nucleic acid, acridine orange may be used as the intercalating dye. The reaction mixtures are transferred into UV transparent plastic tubes spotted with 1% agarose gel, or placed upon UV transparent plastic c and illuminated with 260 nm light. The emission from the DNA-ethidium bromide complex is recorded on film by a camera equipped with an appropriate UV filter. The ability of an agent to condense DNA is inversely proportional to the intensity of the fluorescence in each reaction mixture.

The more precise test is a band shift assay. Briefly, this assay is performed by incubating nucleic acids, either labeled or unlabeled, with various concentrations of candidate condensing agents. Test nucleic acids, conveniently plasmid DNA, and condensing agent are mixed at 1:1 to 1:50 w/w ratios. Following incubation, each sample is loaded on a 1% agarose gel and electrophoresed, the gel is then either stained with ethidium bromide or dried and autoradiographed. DNA condensation is determined by the inability to enter the gel compared to a non-condensed standard. Sufficient condensation is achieved when at least 90% of the DNA fails to enter the gel to any significant degree.

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Condensation can also be measure by directly determining the size of the complex using a light scattering instrument such as the a Coulter N4MD submicron analyzer, for example. Polynucleotides and a condensing agent are incubated at an appropriate ratio, either alone or in the present of 2% PEG-2000 (Fisher Scientific), and 0.6 M NaCl., and then diluted into 3 ml of water. This dilute solution is analyzed by the Coulter counter which will detect particles with a mean size of 0-1,000 nanometers (nm). Condensing agents, such as poly-L-lysine, typically yield particles with a mean diameter of approximately 50-200 nm. See Lee et al., 1996, J. Biol. Chem. 271: 8481-8487.

D. Serum and/or Nuclease Protection Properties

The instant polycationic agents are capable of protecting nucleic acids from degradation in serum or from nucleases, including nucleases present in biological fluids, such as serum, prostate, synovial fluid, etc. One advantage of

this type of protection is that smaller amounts of the desired nucleic acids are needed for efficient administration.

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When present in effective amounts, these polycationic agents can inhibit serum degradation by at least 5 minutes as compared with uncomplexed nucleic acids; more usually, the amount used is sufficient to inhibit degradation by at least 10 minutes; even more usually; the amount used is sufficient to inhibit degradation by at least 30 minutes; even more usually, the amount used is sufficient to inhibit degradation by at least 45 minutes; even more usually, the amount used is sufficient to inhibit degradation by at least 60 minutes; even more usually, the amount used is sufficient to inhibit degradation by at least 90 minutes; and more usually, the amount used is sufficient to inhibit degradation by at least 120 minutes.

Increased serum protection can be measured simply by incubation of the polycation/polynucleotide complex with mouse serum, for example. Preferably, the serum will not be heat inactivated. After incubation, the mixture can be analyzed by gel electrophoresis to determine the quantity of the polynucleotides remaining after incubation.

Alternatively, nucleases can be added to the polycationic agent/nucleic acid complexes. The resulting mixture can be analyzed by gel electrophoresis to determine the amount of degradation. Other biological fluids, such as prostate flud, can also be tested.

E. Mediating Entry of Polynucleotides into a Cell

The polycationic agents can mediate entry of polynucleotides into a cell. Incorporation of polynucleotides into a cell can be measured by either protein expression assays or polynucleotide hybridization techniques, for example.

One method of detecting frequency of incorporation is to include a gene that encodes a marker protein, such as luciferase. Cells that have incorporated the delivered polynucleotides will express the marker protein. The protein can be detected by standard immunoassays, or by biological or enzymatic activity, as in the case of luciferase.

Alternatively, standard hybridization techniques, such as Southern or Northern blots or polymerase chain reaction (PCR) techniques, can be used to detect the presence of the desired polynucleotides.

F. Additional Properties

- To facilitate entry of nucleic acids to the interior of cells, the instant agents can be capable of
 - (a) binding the polynucleotide to the cell surface;
 - (b) cell membrane destabilization;
 - (c) triggering endocytosis;
- 10 (d) endosome buffering capacity;
 - (e) releasing DNA/lipid complexes from endosomes; or
 - (f) nuclear tropism.

Assays for detecting these characteristics are standard and known to those skilled in the art.

15 Physical Properties

The following physical characteristics are factors to consider when determining the composition of the polycationic agents:

- (a) distance between the substituents and the backbone
- (b) the total length of the chain;
- 20 (b) hydrophobicity and/or aromacity;
 - (c) number of hydrogen bonding groups; and
 - (c) charge, including
 - (i) type of charge group, (ii) density of charge and (iii) position.

Other relevant characteristics include structural flexibility. For example, a
helical conformation of the polycationic agent may be preferred for some applications.

Specific dimensions to be considered include

- (a) the distance of phosphate groups in the polynucleotide of interest; and
- (b) the distance of monomer groups in the agents of interest.

Polypeptide Polycationic Agents

The following are examples of useful polypeptides as polycationic agents: polylysine, polyarginine, polyomithine, and protamine. Other examples include histones, protamines, human serum albumin, DNA binding proteins, non-histone chromosomal proteins, coat proteins from DNA viruses, such as \$\phi X174\$, transcriptional factors also contain domains that bind DNA and therefore may be useful as nucleic aid condensing agents. Briefly, transcriptional factors such as C/CEBP, c-jun, c-fos, AP-1, AP-2, AP-3, CPF, Prot-1, Sp-1, Oct-1, Oct-2, CREP, and TFIID contain basic domains that bind DNA sequences.

Organic polycationic agents include: spermine, spermidine, and purtrescine.

The dimensions and of the physical properties of a polycationic agent can be extrapolated from the list above, to construct other polypeptide polycationic agents or to produce synthetic polycationic agents.

15 Synthetic Polycationic Agents

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Synthetic polycationic agents which are useful include, for example, DEAE-dextran, polybrene. Lipofectin®, and lipofectAMINE™ are monomers that form polycationic complexes when combined with polynucleotides.

A preferred group of polycationic agents of the present invention have the following general formula (I):

$$Ta - \left[\begin{array}{ccc} R_1 & R_2 & O \\ | & V & || \\ -N - C - C - C - C \\ \tilde{R}_3 & \end{array} \right] - Tc$$

A preferred subset of these compounds include compounds having formula (I) where R_2 is hydrogen. Even more preferred are polymers comprising at least one natural amino acid. Also preferred are polymers where R_2 and R_3 are both hydrogen, also referred to as poly N-substituted glycines or poly NSGs.

A. Monomers

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The polycationic agent of the invention comprises monomers with the following structure (Π):

Generally, R_1 , R_2 , and R_3 are moieties each with a molecular weight from 1 to 250 daltons. More typically, the molecular weight is no more than 200; even more typically, no more than 175.

Typically, each monomer comprises one hydrogen at R_1 , R_2 , or R_3 . More, typically, either R_1 and R_3 are both hydrogen, the structure of an L-amino acid; or R_3 and R_3 are both hydrogen, the structure of a NSG.

Monomers to be utilized in the polycationic agents can be either positively or negatively charged. Also, neutral substituents can also be utilized.

Degradation sites can be incorporated into the polymer, for example, by including substituents from a natural amino acid when R_1 and R_3 are hydrogen. These monomers can be positively or negatively charged, or neutral.

As a general rule, a basically charged monomer has a pKa value for the side chain of at least 7.5. Positively, or basically, charged monomers include without limitation those containing the following functional groups: amino, guanidino, hydrazido, and amidino. These functional groups can be either aromatic or aliphatic.

Positively charged monomers comprising hydrogen at R₃ and R₄, can be included in the polycationic agent, for example, as a degradation site. Such degradation site may aid in separation of the polycationic agent from the polynucleotide to permit further processing. For an L-amino acid like monomer, useful R₂ substitutents are, for example, from those found in naturally occurring amino acids, such as lysine and arginine. Also, sidechains from amino acid analogues can be used such as ornithine and canaline; or modifications of basic amino acids, such as homoarginine, and modifications of other amino acids such as guanidinovalinate, and aminoethylcysteine. The substitutents found in L-

amino acids can also be incorporated at the R_1 and R_2 positions of the instant polycationic agents.

Naturally occurring amino acids and analogues are designated D-amino acids to indicate the chirality of these molecules. L-amino acids can also incorporated as monomers into the polycationic agents. The substituents of L-amino acids can be, for example, the same as those named for the D-amino acids.

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Preferable monomers include N-substituted glycine monomers. Exemplary N-substitutions include alklphenyl, indolylalkyl, alkoxyphenyl, halophenylalkyl, hydroxyphenylalkyl, as well as the N-substitutions shown below.

Alkylammonium, where preferably $R_1 = H$;

 $R2 = H, CH_3;$

R3, R4, R5 can each be CH3, or CH3CH2; and n = 1-6.

$$\begin{array}{c} R_1 \\ N-C-(CH_2)n-NH_3^+ \\ R_2 \end{array}$$

Aminoalkyl,

where preferably $R_1 = H$;

 $R2 = H, CH_3$;

 $R_3 = CH_3$, CH_3CH_2 ; and

n = 1-6.

$$\begin{array}{c} R_1 \\ N - C - (CH_2)n - NH - NH_2^{+} \\ R_2 & NH_2 \end{array}$$

Guanidinoalkyl,

where preferably $R_1 = H$;

 $R2 = H, CH_{3}$

 $R_3 = CH_3$, CH_3CH_2 ; and

n = 1-6.

$$\begin{array}{c|c}
R_1 \\
N-C \\
-(CH_2)n \\
NH_2
\end{array}$$

$$\begin{array}{c}
NH_2^+ \\
NH_2
\end{array}$$

Amidinoalkyl,

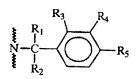
where preferably $R_1 = H$:

 $R2 = H, CH_3;$

 $R_3 = CH_3$, CH_3CH_2 ; and

n = 1-6.

Aminocyclohexyl



Guanidinobenzyl

where preferably $R_1 = H$; and

 $R_2 = H, CH_3$

R3, R4, and R5 each can be H or _

Piperidyl

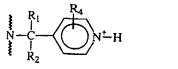
$$R_3$$
 R_4
 R_5
 R_5

Amidinobenzyl

where preferably $R_1 = H$;

 $R_2 = H$, CH_3 ; and

R3, R4, R5 each can be.



Pyridylmethyl

where preferably

 $R_1 = H;$

 $R_2 = H$, CH_3 ; and

 $R_4 = H, CH_3O, Cl, F, Br, CH, NO_2, CH_3.$



Aminobenzyl where preferably

 $R_1 = H$;

 $R_2 = H$, CH_3 ; and

 $R_4 = H$, CH_3O , CI, F, Br, CH. NO_2 , CH_3 .

The positively charged substituents described above can also be placed at the R, or R, positions of formulas (I) and (II).

The polycationic agents can comprise negatively charged or neutral monomers. As with the positively charged monomers, D-amino acid, L-amino acid, and NSGs are preferred to be incorporated as monomers.

The following are examples of such monomers:

B. <u>Polycationic Polymers</u>

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Typically, the polycationic agents exhibit a predicted isoelectric point of at least 9, excluding the terminal groups. Further, the agents contain, excluding the terminal groups, at least 20% positively charged monomers; more typically, at least 25% more typically, 30%; and preferably, at least 33% positively charged monomers. Typically, the agents do not comprises greater than 5% acidic monomers and preferably none.

The charge density and composition of the polycationic agent can be altered to accommodate the specific nucleic acid sequence, type, and other components included with the complex of nucleic acids and polycationic agent.

Usually, the length of the polymer is at least 8 monomers; even more usually, 12 monomers; even more usually, 18 monomers. More typically, the polycationic agents of the invention will be at least 24 monomer units in length; more typically, 30 monomer units; even more typically, 36 monomer units; even more typically, 48 monomer units. The polycationic agent can be up to 50 to 75 to 100 monomer units in length.

Preferably, the polycationic agent comprises monomers where all R_2 and R_3 are hydrogen. Even more preferably, where all R_2 and R_3 are hydrogen, the polycationic agent comprise repeating trimer units with the following monomer sequence (from amino to carboxy terminus): (1) neutral monomer, (2) neutral monomer, and (3) positively charged monomer.

Preferably, the neutral monomer comprises an aromatic group at the R_1 position; more preferably, wherein the aromatic group comprises a single ring; even more preferably, wherein the aromatic group is a six member ring.

Typically, the positively charged monomer is aminoalkyl at the R_1 position; more typically, the aminoalkyl comprises 1-6 carbon molecules; even more typically, the aminoalkyl is aminoethyl.

Typically, the polycationic agent comprises between 3 to 20 repeating trimers, trimers having two neutral and one positively charged R₁ groups are preferred, such as, for example, trimer shaving the sequence, neutral monomer, neutral monomer, positively charged monomer. More preferably, the polycationic agent comprises 5 to 18 trimers; preferably 8 to 16 trimers; and even more preferably, 12 to 16 trimers.

Optionally, the polycationic agent contains only positively charged monomers, excluding the terminal groups. Typically, such a polycationic agent comprises between 24 and 48 monomers; more typically, 30 to 40 monomers; even more typically, 36 monomers.

Polycationic agents of the present invention containing only positively charged monomers typically have guanidinoalkyl sidechains. Typically, the guanidinoalkyl sidechain comprises 1 to 6 carbon molecules. Preferably, the side chain is guanidino ethyl.

C. Neutral Polymers

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A preferred group of neutral polymers of the present invention have the general formula (I):

$$Ta - \begin{bmatrix} R_1 & R_2 & O \\ -N - C - C - C \\ R_3 & T \end{bmatrix} - Tc$$

Preferably, R_2 is hydrogen. Even more preferred are polymers comprising at least one natural amino acid. Also preferred are polymers having formula (I) where R_2 and R_3 are hydrogen, also referred to as poly N-substituted glycines or poly NSGs.

Monomers employed in neutral polymers of the present invention have the same general formula as monomers employed in cationic polymers of the present invention, i.e.:

Generally, R₁, R₂, and R₃ are moieties each with a molecular weight from 1 to 250 daltons. More typically, the molecular weight is no more than 200; even more typically, no more than 175.

Typically, each monomer comprises one hydrogen at R_1 , R_2 , or R_3 . More, typically, either R_1 and R_3 are both hydrogen, the structure of a L-amino acid; or R_2 and R_3 are both hydrogen, the structure of a NSG.

Monomers to be utilized in the neutral agents can be either positively or negatively charged. Also, neutral substituents can also be utilized. Neutral polymers exhibit no net positive or negative charge, excluding the terminal groups.

Degradation sites can be incorporated into the polymers by using naturally occurring amino acid substituents in monomers when R_1 and R_2 are hydrogen.

Naturally occurring amino acids and analogues are designated D-amino acids to indicate the chirality of these molecules. L-amino acids can also incorporated as monomers into the neutral polymers. The substituents of L-amino acids can be, for example, the same as those named for the D-amino acids.

Preferred monomers include N-substituted glycine monomers, and monomers that are capable of forming hydrogen bonds and/or ionic bonds with the polynucleotides to be delivered.

Examples of monomers for the neutral polymers include those described above and in the Examples below.

D. Linking Polymers Together

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Polymers can be linked together incorporating terminating groups or sidechains that permit cross-linking of the polymers. For example, polymers can be linked by a disulfide bond. Other terminating groups useful for coupling polymers include, carbonate, urea, and the like.

E. Additional Groups to be Incorporated into the Polymer

Additional components can be included in the polycationic agents of the instant invention, such as targeting ligands. Such additional groups can facilitate endocytosis of the desired nucleic acids or aid binding of the nucleic acids to the cell surface.

Polypeptides can be incorporated into the polycationic agents. Examples include, without limitation: asialoorosomucoid (ASOR); transferrin; asialoglycoproteins; antibodies; antibody fragments; ferritin; interleukins; interferons, granulocyte, macrophage colony stimulating factor (GM-CSF), granulocyte colony stimulating factor (G-CSF), macrophage colony stimulating

factor (M-CSF), stem cell factor and erythropoietin. Viral antigens, such as envelope proteins, can also be used. Also, proteins from other invasive organisms are useful, such as the 17 amino acid peptide from the circumsporozoite protein of plasmodium falciparum known as RII.

In addition, lipoproteins can be incorporated into the polycationic agent, such as low density lipoprotein, high density lipoprotein, or very low density lipoprotein. Mutants, fragments, or fusions of these proteins can also be used.

Other groups that can be incorporated include without limitation: hormones, steroids, androgens, estrogens, thyroid hormone, or vitamins, folic acid. Folic acid can be incorporated into the polycationic agent according, for example, to Mislick et al., 1995, T.J. Bioconjugate Chem. 6:512.

Also, the polycationic agents of the instant invention can be chemically conjugated with polyalkylene glycol. In a preferred embodiment, the polyalkylene glycol is polyethlylene glycol. PEG can be incorporated with a polycation agent according, for example, to Lu et al., 1994, Int. J. Pept. Protein Res. 43:127.

In addition, the polycationic agent can be chemically conjugated with mono-, di-, or polysaccaride. In a preferred embodiment of this aspect, the polysaccharide is dextran.

These additional groups can be incorporated within the polycationic agent. For example, R_1 , R_2 , and R_3 can be a substituent that is capable of being activated to cross link with any one of the above groups. For example, a thiol group could be included to cross link with another group to form a disulfide bond.

F. <u>Terminal Groups</u>

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The terminal groups of the instant polycationic agents can be chosen as convenient. Suitable terminal groups (i.e., Ta and Tc) include, for example, -NH₂, -OH, -SH, and -COOH. Terminal groups can be selected to enhance the targeting properties of the polycationic agent and can be any of the additional groups described above.

The additional groups described above can be incorporated at the terminus of the polycationic agent. For example, the polycationic agent can be

(1) acylated with a variety of carboxylic acids; (2) sulfonylated with sulfonyl chlorides; or (3) derivatized with isocyanates or isothiocyanates. Once activated, the terminus can be reacted with any of the above-mentioned groups, such as a polypeptide, such as low density lipoprotein, or folic acid.

One means of adding a terminal group to the polycationic agent is, for example, is (1) to acylate the amino terminus with Fmoc-amino-hexanoic acid; and (2) to remove the protecting group, Fmoc, to generate a primary amine, which can be further functionalized.

Alternatively, the amino-terminal groups can include, without limitation: acyl, such as acetyl, benzoyl; or sulfonyl, such as dansyl.

Carboxy terminal groups can include, for example, amide or alkyl amide.

Synthesis of Polycationic Agents

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Polycationic agents of the present invention can be synthesized by either solid or solution phase methods. The following is a solid phase method for the synthesis of NSGs, which can be generally used for a wide variety of side-chain substitutents. This method can be performed utilizing automated peptide synthesis instrumentation to permit rapid synthesis of polycationic agents of interest. Such instruments are commercially available from, for example, Applied Biosystems and Milligen.

A. Two Step Monomer Assembly

A method of synthesis is to assemble the monomer from two submonomers in the course of extending a polymer comprising an NSG monomer. This technique is described in Zuckermann et al., 1992, J Amer Chem Soc 114(26):10646-10647, and Zuckermann et al., PCT Patent Publication No. WO 94/06451. The NSGs can also be considered to be an alternating condensation of copolymer of an acylating agent and an amine.

The direction of polymer synthesis with the submonomers occurs in the carboxy to amino direction. The solid-phase assembly for each monomer, in the course of polymer formation, eliminates the need for $N\alpha$ -protected monomers, as only reactive side-chain functionalities need to be protected.

Each monomer addition comprises two steps, an acylation step and a nucleophilic displacement step as shown in Figure 1.

Specifically, each cycle of monomer addition consists of two steps:

- (1) acylation of a secondary amine bound to the support with an acylating agent comprising a leaving group capable of nucleophilic displacement by an amine and a carbonyl group, preferably carboxyl. An example is a haloacetic acid; and
- (2) nucleophilic displacement of the leaving group with a sufficient amount of a submonomer comprising a primary amino group to introduce a sidechain. The amino group containing submonomer can be an alkoxyamine, semicarbazide, acyl hydrazide, substituted hydrazine or the like.

Acylation can be activitated with carbodiimide or other suitable carboxylate activation method.

The efficiency of the displacement is modulated by the choice of halide, e.g., I>Cl. Protection of aliphatic hydroxyl groups, carboxylic acids, carboxy, thiol, amino, some heterocycles, and other reactive side-chain functionalities is preferred to minimize undesired side reactions. However, the mild reactivity of some side-chain moieties toward displacement or acylation may allow their use without protection., e.g., indole, imidazole, and phenol.

B. Three Step Monomer Assembly

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NSGs can also be constructed utilizing a three step method for assembling each monomer as the polymer is extended. The backbone of the monomer of first extended by acylation step followed by a nucleophilic displacement. The side chain is introduced by a second acylation step. The reaction scheme is shown in Figure 2.

The backbone of the monomer is assembled in the first two steps of the synthesis cycle. The first reaction is an acylation step where the carbonyl group of the acylating agent reacts with an amine. The acylating agent comprises a carbonyl group; a backbone, R_i; and a leaving group, L. Preferably, the carbonyl group is carboxyl.

The second step is a nucleophilic displacement of the leaving group by the first amino group of the displacing agent. The displacing agent comprises a

first and a second amino group and a backbone, R_d . The first amino group is a primary amine, and the second step produces a secondary amine.

The third step is another acylation in which the another acylating submonomer reacts with the first amino group of the displacing agent to produce a tertiary amide. The acylation agent comprises a carbonyl group; an optional linker; and a sidechain. Preferably, the carbonyl group is carboxyl.

Pharmaceutical Compositions

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The polycationic agent/polynucleotide complexes, whether or not encapsulated in liposomes, may be administered in pharmaceutical compositions. The pharmaceutical compositions comprise a therapeutically effective amount of nucleic acids.

The term "therapeutically effective amount" as used herein refers to an amount of a therapeutic agent sufficient to detectably treat, ameliorate, or prevent a particular disease or condition, i.e., an amount sufficient to induce a detectable therapeutic or preventative effect. The effect may include, for example, chemical markers or antigen levels. Therapeutic effects also include reduction in physical symptoms, such as decreased body temperature. The precise effective amount for a subject will depend upon the subject's size and health, the nature and extent of the cardiovascular condition, and the therapeutics or combination of therapeutics selected for administration. Thus, it is not useful to specify an exact effective amount in advance. However, the effective amount for a given situation can be determined by routine experimentation and is within the judgment of the clinician. For purposes of the present invention, an effective dose will be from about 0.01 mg/kg to 50 mg/kg or 0.05 mg/kg to about 10 mg/kg of the DNA constructs in the individual to which it is administered.

A pharmaceutical composition can also contain a pharmaceutically acceptable carrier. The term "pharmaceutically acceptable carrier" refers to a carrier for administration of a therapeutic agent, such as antibodies or a polypeptide, genes, and other therapeutic agents. The term refers to any pharmaceutical carrier that does not itself induce the production of antibodies harmful to the individual receiving the composition, and which may be administered without undue toxicity. Suitable carriers may be large, slowly

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metabolized macromolecules such as proteins, polysaccharides, polylactic acids, polyglycolic acids, polymeric amino acids, amino acid copolymers, and inactive virus particles. Such carriers are well known to those of ordinary skill in the art.

Pharmaceutically acceptable salts can be used therein, for example, mineral acid salts such as hydrochlorides, hydrobromides, phosphates, sulfates, and the like; and the salts of organic acids such as acetates, propionates, malonates, benzoates, and the like. A thorough discussion of pharmaceutically acceptable excipients is available in *Remington's Pharmaceutical Sciences* (Mack Pub. Co., N.J. 1991).

Pharmaceutically acceptable carriers in therapeutic compositions may contain liquids such as water, saline, glycerol and ethanol. Additionally, auxiliary substances, such as wetting or emulsifying agents, pH buffering substances, and the like, may be present in such vehicles. Typically, the therapeutic compositions are prepared as injectables, either as liquid solutions or suspensions; solid forms suitable for solution in, or suspension in, liquid vehicles prior to injection may also be prepared. Liposomes are included within the definition of a pharmaceutically acceptable carrier.

Delivery Methods

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Once formulated, the compositions of the invention can be administered (1) directly to the subject; (2) delivered *ex vivo*, to cells derived from the subject; or (3) *in vitro* for expression of recombinant proteins. The subjects to be treated can be mammals or birds. Also, human subjects can be treated.

Direct delivery of the compositions will generally be accomplished by injection, either subcutaneously, intraperitoneally, intravenously or intramuscularly or delivered to the interstitial space of a tissue. The compositions can also be administered into a tumor or lesion. Other modes of administration include oral and pulmonary administration, suppositories, and transdermal applications, needles, and gene guns or hyposprays. Dosage treatment may be a single dose schedule or a multiple dose schedule.

Methods for the ex vivo delivery and reimplantation of transformed cells into a subject are known in the art and described in e.g., International Publication No. WO 93/14778 (published 5 August 1993). Examples of cells useful in ex

vivo applications include, for example, stem cells, particularly hematopoetic, lymph cells, macrophages, dendritic cells, or tumor cells.

Generally, delivery of nucleic acids for both ex vivo and in vitro applications can be accomplished by the following procedures, for example, dextran-mediated transfection, calcium phosphate precipitation, polybrene mediated transfection, protoplast fusion, electroporation, encapsulation of the polynucleotide(s) in liposomes, and direct microinjection of the DNA into nuclei, all well known in the art.

The examples presented below are provided as a further guide to the practitioner of ordinary skill in the art, and are not to be construed as limiting the invention in any way.

Example 1

Synthesis of Polycationic Agents

This example describes the synthesis of polycationic agents with the following structure:

$$Ta - \begin{bmatrix} -R_1 & R_2 & O \\ -N - C - C - C \end{bmatrix}_{n} - Tc$$

where R_3 and R_2 are hydrogen for all monomers. All polymers describe in this example terminate in an amino and a carboxyl group unless specified, such as a folate terminating group.

The polycationic agents described below were synthesized according to the procedures described in Figliozzi et al., 1996, Meth. Enzy. 267:437-447, and Zuckermann et al., 1992, J. Amer. Chem. Soc. 114(26):10646-10647.

All polymers were synthesized using bromoacetic acid and primary amines. The following are substitutents of the primary amines to be positioned at R_1 to construct the polycationic agents:

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Cationic Sidechains

$$p = (S)-1$$
-methylethylenediamine

Other Sidechains

Nm = naphthylmethyl (R = H)
$$5N = (S) - \alpha - methylmaphthylmethyl (R = CH_2)$$

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Other Sidechains

Peptoid-Folic acid conjugates

- Fmoo-aminohexanolc acid
 2. 20% piperidine/DMF

- 0.1 M Folic add in DMSO, DIC HOBt, 50°
 Triffuoroacetic add

Abbreviation	Description
Bn	benzyl
Chm	cyclohexylmethyl
Ff	furfurylmethyl
G	guanidinoethyl
Gp	guanidinopropyl
Н	
H+	(S) alpha-methylbenzyl
Me	(R) alpha-methylbenzyl
Nm	methoxyethyl
P	naphthylmethyl
P,	aminoethyl
P*	aminohexyl
	(S)-α-methylaminoethyl
Ph	phenethyl
Pr	aminopropyl
Py	N-pyrrolidinopropyl
Tmb	3,4,5,-trimethoxybenzyl
Q	trimethylaminoethyl
Phpr	phenylpropyl
6-gal	6-galactosyl
Trp	N-2-(3-indolylethyl)
pMeOph	p-methoxyphenethyl
pClPh	p-chlorophenethyl
Tyr	p-hydroxyphenethyl
sN	(S)-α-methylnaphthylmethyl

The polycationic agents synthesized include:

Name	Sequence	Length	Mol. Wt.	# charges
RZ110-1	(HHP)6	18	2550.8	7
RZ110-2	(HP)9	18	2367.8	10
RZ110-3	(HPP)6	18	2184.8	13
RZ110-4	(HPPP)4HP	18	2123.8	14
RZ110-5	(HHP')6	18	2869.8	17
RZ110-6	(HP')9	18	2871.8	10
RZ110-7	(HP'P')6	18	2856.8	13
RZ110-8	(HP'P'P')4HP'	18	2851.8	14
RZ110-9	(HHP)12	36	5084.6	13
RZ110-10	(HP)18	36	4718.6	19
RZ110-11	(HPP)12	36	4352.6	25
RZ110-12	PP(HPPP)8HP	36	4230.6	27
RZ110-13	(HHP')12	36	5722.6	13
RZ110-14	(HP')18	36	5726.6	19
RZ110-15	(HP'P')12	36	5696.6	25

Name	Sequence	Length	Mol. Wt.	# charges
RZ110-16	5 P'P'(HP'P'P')8HP'	36	5686.6	27
RZ112-1	(Q)36	36	5181.5	37
RZ112-2	(G)36	36	5130.8	37
RZ112-3	(HP*P*P*)9	36	4529.3	28
RZ112-4	(P*)36	36	4122.8	37
RZ112-5	(HP*P*P*)4HP*	18	2305.2	14
RZ112-6	(P*)18	18	2069.9	19
RZ112-7	(P)18	18	1817.7	
RZ112-8	(P)36	36	3618.4	19
RZ120-1	(MeMeP)8	24	2658.4	37
RZ120-2	(BnBnP)8	24	3170.8	9
RZ120-3	(HHP)8	24	3394.8	9
RZ120-4	(H+H+P)8	24	3394.8	9
RZ120-5	(MeMeP)12	36	3979.2	
RZ120-6	(BnBnP)12	36	· 	13
RZ120-7	(HHP)12	36	4747.6	13
RZ120-8	(H+H+P)12	36	5083.6	13
RZ120-9	(MeMeP)16	48	5083.6 5299.9	13
RZ120-10	(BnBnP)16	48	6324.5	17
RZ120-11	(HHP)16	48	6772.5	17
RZ120-12	(H+H+P)16	48	6772.5	17
RZ120-13	(HHP)12 folate	36	5300	17.
RZ123-1	(HHPr)12	36	5252	13
RZ123-2	(HHPr)12	36	5252	13
RZ123-3	(HHP)12	36	5084	13
RZ123-4	folate-(HHPr)12	36		13
RZ123-5	(HHGp)12	36	5862 5756	13
RZ123-6	(HHG)12	36	5588	13
RZ124-1	(HHPr)12	36	5252	13
RZ124-2	(sNsNPr)12	36	6452	13
RZ124-3	(NmNmPr)12	36	6116	13
RZ124-4	(PyPyPr)	36	5756	13
RZ124-5	(HHPy)12	36	6069	13
RZ124-6	(Py)36	36	6573	13
RZ124-7	folate-(HHPr)12 -	36	5862	
RZ127-1	(PhPhP)12		5085	13
RZ127-2	(ChmChmP)12		4895	13
RZ127-3	(TmbTmbP)12		6912	13
RZ127-4	(FfFfP)12			13
RZ136-3	(PhprPhprP)12		4508	13
RZ140-2	(6-gal)12-(PhPhP)12		5419	13
RZ140-3	(TrpTrpP)12		7712	13
RZ144-1	(PhPPh)12		6020	13
RZ144-2	(PPhPh)12		5083	13
RZ144-3	(pMeoPhpMeoPhP)		5083	13
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Sequence	Length	Mol. Wt.	# charges
(pClPhpClPhP)12	36		13
AMCA-(PhPhP)12	36		12
(TyrTyrP)12			12
(6gal6galP)12			13
	~~~ <del> </del>		
			13
	(pClPhpClPhP)12 AMCA-(PhPhP)12	(pClPhpClPhP)12 36 AMCA-(PhPhP)12 36 (TyrTyrP)12 36 - (6gal6galP)12 36 (PhPhP)12 36	Wt.   Wt.   Wt.     Wt.     Wt.

*purified

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To summarize the method, Fmoc-Rink amide resin (NovaBiochem, San Diego, California, USA) is used as the solid support. This is the same resin that is used for the Fmoc synthesis of peptide C-terminal amides. The polycationic synthesis begins with the deprotection of the Fmoc group on the resin with 20% (v/v) piperidine-dimethylformamide (DMF). The amino resin is then acylated with bromoacetic acid. This is followed by nucleophilic displacement of the bromide with a primary amine to build the NSG monomer. The latter two steps are then continued in an iterative fashion to elaborate the desired oligomer.

All reactions and washings were performed at room temperature unless otherwise noted. Washing of the resin refers to the addition of a wash solvent (usually DMF or dimethylsulfoxide (DMSO)) to the resin, agitating the resin so that a uniform slurry is obtained (typically for about 20 seconds), followed by thorough draining of the solvent from the resin. Solvents were removed by vacuum filtration through the fritted bottom of the reaction vessel until the resin appeared dry (typically about 5 seconds). In all the syntheses, resin slurries were agitated via bubbling argon up through the bottom of the fritted vessel.

A fritted reaction vessel was charged with 100 mg (50  $\mu$ mol) of Frnoc-Rink amide resin with a substitution level ~ 0.50 mml/g resin. Two milliliters of DMF was added to the resin and this solution was agitated for 1-2 minutes to swell the resin. The DMF was then drained. The Frnoc group was then removed by adding 2.0 ml of 20% piperidine in DMF to the resin. This was agitated for 1 minute and then drained. Another 2 ml of 20% piperidine in DMF was added to the resin and agitated for 15 minutes and then drained. The resin was then washed with DMF, six times with 2 ml.

The deblocked amine was then acylated by adding 850 $\mu$ l of 0.6 M bromoacetic acid in DMF to the resin followed by 200  $\mu$ l of 3.2 M N,N'-

diisoprooplycarbodiimide (DIC) in DMF. This solution was agitated for 30 minutes at room temperature and then drained. This step was repeated a second time. The resin was then washed with DMF, twice with 2 ml and DMSO, once with 2 ml. This completed one reaction cycle.

The second cycle was initiated by the acylating rep with bromoacetic acid and DIC, followed by displacement with the second amine. This acylation/displacement cycle was repeated until the desired oligomer was obtained.

Cleavage of the resin from the polycationic agent is as follows. The dried resin was placed in a glass scintillation vial containing a teflon-coated micro stir bar, and approximately 5 ml of 95% trifluoroacetic acid (TFA) in water was added. The solution was stirred for 20 minutes and then filtered through an 8-ml solid-phase extraction (SPE) column fitted with a 20-µm polyethylene frit into a 50 ml polypropylene conical centrifuge tube.

The resin was washed with 1 ml 95% TFA. The combined filtrates were then lyophilized three times from 1:1 acetonitrile:water. Material was redissolved to a concentration of 5 mM in 5% acetonitrile in water.

## Preparation of guanidinoalkyl-containing Polymers:

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The guanidinoalkyl sidechains were introduced into the polymers by post-synthesis modification of aminoalkyl sidechains. Thus, polymers were synthesized by the submonomer method as described above except that methoxybenzhydrylamine (MBHA) resin was used instead of the Rink resin. Wherever a guanidinoalky sidechain was desired, a mono-Boc-alkanediamine was incorporated in the displacement step. After elaboration of the polymers, the sidechain Boc groups were removed by treatment with 95% TFA/water for 20 min at room temp. (This does not remove the oligomer from the soliud support). The free amino groups were then guanidinylated by treatment with 1 H-pyrazole-1-carboxamidine (1 M in DMF, 2 x 1 hr, 40°C). After washing with DMF and methylene chloride, the oligomer was cleaved from the resin with hydrofluoric acid, and lyophilized.

## Preparation of folic acid - polymer conjugates:

Folic acid - polymer conjugates were prepared by adding a linker to the N-terminus of the resin-bound polymer which was then acylated with folic acid. Specifically, after elaboration of the polymer, the N-terminus was acylated with Fmoc-aminohexanoic acid (0.5 M in DMF, 0.5 M hydroxybenzotriazole, 0.5 M diisopropylcarbodiimide (DIC), 1 x 1 hr, room temp.). After Fmoc group removal (20% piperidine/DMF, 1 x 20 min, room temp.), the free primary amino group was acylated with folic acid (0.1 M in DMSO, 0.1 M DIC, 1 x 2 hr, 50°C). After washing of the resin, the conjugate was cleaved with 95% TFA/water in the usual fashion.

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#### Example 2

## Condensation of Polynucleotides

Polycationic agents were synthesized and isolated to a final concentration of 5 mM as described in Example 1. Polynucleotides were condensed with RZ110, RZ112, and RZ120 series compounds according to the following procedure.

- (1) Dilute all polycationic agents to a final concentration of 3 nanomoles of positive charge per microliter.
  - (2) Add 1  $\mu$ g of DNA to 1-2 $\mu$ l of diluted polycationic agents.
- (3) Adjust volume to 10  $\mu$ l. This mixture can be stored overnight at 20 4°C.
  - (4) Add of 5  $\mu$ l of DNA/polycationic mixture to 2  $\mu$ l of 5X buffer, which does not contain SDS to maintain the complex. (5X buffer = 40% sucrose, 0.25% bromphenol blue and 200 mM Tris Acetate, 4 mM EDTA (PH 7.8).
    - (5) Adjust volume to 10 μl.
  - (6) Run sample on a 1% agarose gel utilizing 75 volts for 1.5 hours.. Between 1 to 2 μl, all polycationic agents were judged to retard the migration of DNA into an agarose gel.

#### Example 3

### Inhibition of Serum Degradation

The RZ110, RZ112, and RZ120 series compounds were mixed with polynucleotide as described in Example 2. Five microliters of the overnight mixture was added to 5 µl of BalbC mouse serum. The serum was not heat treated but freeze thawed. The serum, polycationic agent, and polynucleotide mixture was incubated typically for 30 minutes at 37°C. The time of incubation varied between 5 and 60 minutes

Next, 2  $\mu$ l of 5X buffer containing 0.5% (wt/v) SDS was added to the incubated mixture. This final solution was loaded onto a 1% agarose gel and electrophoresed at 75 volts for 1.5 hours.

All of the compounds tested, i.e., the entire RZ110, 112, and 120 series, provided significant protection in a direct comparison. The entire RZ112 series and RZ110-3 and RZ110-8 inhibited serum degradation better than poly-L-lysine.

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#### Example 4

#### Peptoid Mediated In Vitro Delivery

DNA comprising a luciferase gene  $l\mu g/\mu l$ , was diluted into endotoxin free water. The plasmid DNA was CMVKm luciferase, which is described in more detail in Example 5.

The transfection protocol for in vitro delivery was as follows:

- (A) HT1080 cells were used. These cells are available from American Type Culture Collection, Rockville, Maryland, USA, Accession No. CCL 121. This is a fibrosarcoma. The growth medium was Dulbecco's Modified Eagle medium (DME) with 10% heat-inactivated fetal calf serum.
- 25 (B) Twenty four hours prior to transfection, the cells were placed at 5 X 10⁴ per well of a 24-well plate in 1 ml of medium.
  - Feed cells with 500μl of DME-10% fetal calf serum (FCS )or 500μl
     Opti-MEM®. Opti-Mem® can be purchased from Gibco BRL, Life
     Technologies, Inc., Gaithersburg, Maryland, USA.
    - 2. Add 200 μl Opti-MEM® to each tube.

- Add 3  $\mu l$  of the desired polycationic agent to the 200 $\mu l$  of Opti-3. MEM®.
  - Add 2 µl of 1µg/µl luciferase DNA, mix. 4.
  - Incubate mixture for 5 minutes at room temperature. 5.
- Add 100 µl of the polycationic agent/DNA mixture to plate with 5 6. DME-FCS, 100  $\mu$ l to cells fed with Opti-MEM®.
  - Incubate cells and polycationic agent/DNA mixture for ~4 hours at 7. 37°C.
    - 8. Change media on all cells to DME-FCS.
- DME-FCS was used as a positive control. As a control, a transfectant, LT1, was used from Panvera, Inc., Madison, Wisconsin, USA to transfect cells in serum and cells in Opti-MEM®.
- 10. Cells were tested for luciferase activity using a Promega Luciferase Assay System from Promega, Madison, Wisconsin, USA., in accordance with 15 the manufacturer's directions.

#### Results:

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Name	Formula	Luciferase (RLU)
RZ120-1	(MeMeP)8	0
RZ120-2	(BnBnP)8	0.93
RZ120-3	(HHP)8	1.38
RZ120-4	(H+H+P)8	1.5
RZ120-5	(MeMeP)12	0
RZ120-6	(BnBnP)12	1.64
RZ120-7	(HHP)12	2.64
RZ120-8	(H+H+P)12	2.84
RZ120-9	(MeMeP)16	0
RZ120-10	(BnBnP)16	1.42
RZ120-11	(HHP)16	1.94
RZ120-12	(H+H+P)16	1.32
control	LT1	51.96

Experiment #2		
Name	Formula	Luciferase
		(RLU)
RZ110-1	(HHP)6	0.0015
RZ110-2	(HP)9	0.0012
RZ110-4	(HPPP)4HP	0.0004
RZ110-5	(HHP')6	0.0006
RZ110-6	(HP')9	0.0052
RZ110-7	(HP'P')6	0.0005
RZ110-8	(HP'P'P')4HP'	0.0003
RZ110-9	(HHP)12	8.7
RZ110-10	(HP)18	0.0014
RZ110-12	PP(HPPP)8HP	0.0459
RZ110-13	(HHP')12	2.5
RZ110-14	(HP')18	2.2
RZ110-15	(HP'P')12 '	0.064
RZ110-16	P'P'(HP'P'P')8	0.01
control	LT I	88.7

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#### Example 5

#### Targeting Ligand

A. Cells, Vector, and Compositions Used.

In a first experiment, murine endothelial cells (Py-4-1) which express high levels of acetylated-LDL receptors. The cells and the LDL receptors are described in Dubois et al., 1991, Exp. Cell Res. 196:302-313.

A luciferase-containing plasmid (pCMVkmLUC) was used to determine if polynucleotides could be delivered and expressed into endothelial cells when associated with polycationic agents described in Example 1 with acetylated-LDL (Ac-LDL). A description of the identification and isolation of endothelial cells based on their increased uptake of acetylated-low density lipoprotein is in Voyta et al., 1984, J. Cell Biol. 99: 2034-2040.

The plasmid used in these experiments pCMVkmLUC, was constructed by inserting the luc + gene from pSP-luc+ (Promega Corporation, Madison, WI) into the expression vector pCMVkm2. Briefly, pSP-luc+ was digested with the restriction enzymes Nhe1-EcoRV (Boehringer Mannheim, Indianapolis, IN) and a fragment of 1691 bp was isolated by standard methods. This fragment was inserted into pCMVkm2, which had been digested with XbaI and EcoRV using the Rapid Ligation Kit (Boehringer Mannheim, Indianapolis, IN). The sequence of pCMVkm2 is depicted in SEQ ID NO:2 and described below. The luc + gene was cloned into pCMVkm2 such that expression is driven by the CMV immediate early enhancer promoter and terminated by the bovine growth hormone polyadenylation signal.

The luciferase expression was compared to levels obtained with the same vector delivered in conjunction with lipofectamine, an agent used commonly to transfect cells *in vitro* (Hawley-Nelson et al., 1993, Focus 15:73). The results are presented in the table below.

#### B. Method of Transfection:

Briefly, the cells were plated in 24 well dishes, grown to approximately 80% confluence, transfected and assayed 24 hours later for luciferase activity. All transfections were done in serum containing medium. During transfection mixture preparation, pCMVkmLUC was first mixed with RZ 112, and the DNA-

cationic polycationic agent complexes were then added to Ac-LDL. Serum containing medium was then added to the mixtures to adjust the volume delivered to each well to 0.5 ml.

Lipofectamine was used as a positive control. No lipoprotein was added to this positive control. Lipofectamine is a 3:1 (w/w) liposome formulation of the polycationic lipid 2,3,-dioleylosy-N-[2(spermine-carboxamido)ethyl]-N,N-dimethyl-1-propanaminiumtrifluoroacetate (DOSPA) and the neutral lipid dioleoyl phosphatidyl-ethanolamine (DOPE) in membrane-filtered water. Lipofectamine can be purchased from Life Technologies, Gaithersburg, Maryland, USA).

#### C. <u>Luciferase Assay</u>

Luciferase activity was assayed using the Promega Luciferase Assay System, Madison, Wisconsin.

#### 15 D. Results

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Table 1 shows the results of an experiment where the polycationic agent, RZ-112-2 was compared to lipofectamine to deliver the luciferase gene to cells comprising the acetylated LDL receptor.

Table 1
LuciferaseActivity

Group	Ac-LDL (μg)	pCMVkmLUC (µg)	RZ 112-2 (nm)	pg luc/mg
1	5	10	2.5	61
2	-	10	-	0
4	0.5	10	2.5	16
5	•	1	0.25 . 0.25	16 17
6	0.5	10	2.5	631
/	0.5	10	5	1996
IPOFEC	TAMINE CO	NTROL		
}	<u>.</u>	10	-	10786

^{*} each number represents the mean of three wells.

#### Example 6

## Comparison of Cells with and without Acetylated LDL Receptors

### A. Cells with Acetylated LDL Receptors

For this experiment, K1735 mouse, epithelial melanoma cells were used.

These cells express low or non-existent levels of Ac-LDL receptors. A description of the cells is in J. Natl. Cancer Inst. 69(4): (1982).

#### B. Methods

Briefly, the cells were plated in 24 well dishes at 10,000 cells per well in DME with 10% FCS supplemented with 2 mM L-glutamine. The Py-4-1 cells were cultured in 10% CO₂ at 37°C. The K1735 cells were cultured in 5% CO₂ at 37°C. The cells were grown to approximately 50% confluence, transfected and assayed 24 hours later for luciferase activity. All transfections were done in serum containing medium.

During transfection mixture preparation, pCMVkmLUC was first mixed with RZ 112-2, and the DNA-polycationic agent complexes were then added to Ac-LDL. Serum containing medium was then added to the mixtures to adjust the volume delivered to each well to 0.5 ml.

#### C. Results

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<u>Table 2</u> <u>Luciferase Activity</u>

Group	Ac-LDL (µg)	pCMVkmLUC (µg)	RZ 112-2 (nm)	pg luc/m	g protein*
		<b>""</b> 37	, ,	Py-4-1	K1735
1	0.5	1	1	1301	24
2	0.5	i	5	2181	0
3	0.5	1	10	373	0
4	0.5	10	5	840	0
5	-	1	5	327	0
6	-	1	5	945	ND
10	5	1	5	298	ND
LIPOFE	CTAMINE	CONTROL			112
7	-	1	<u> </u>	23	0
8	-	10	<del>-</del>	2878	960

^{*} each number represents the mean of three wells.

# Example 7 Injection of Polynucleotides Encoding Erythropoietin

#### A. Polynucleotides

CMVkm2 is the standard vector used in these studies. CMVkm2 is a vector optimized for expression in mammalian cells. The gene of interest is cloned into a polylinker which is inserted 3' of a human CMV expression cassette. This cassette contains the human CMV immediate early promoter/enhancer followed by intron A of the human CMV immediate early region (Chapman et al., 1991, Nucl. Acids Res. 19:3937-3986). Transcription is terminated by a polyadenylation site from the bovine growth hormone gene, which has been cloned immediately 3' of the polylinker. See SEQ ID NO:2 for the CMVkm2 vector.

The CMV-km-cmEPO vector was constructed from CMVkm2 as follows. The cynomolgus monkey EPO cDNA was acquired from the ATCC (Accession No. 67545, Rockville, MD). This plasmid was cut with <u>AvrII</u> and <u>BglII</u> and inserted into the <u>XbaI</u> and <u>BamHI</u> sites of the CMVkm2 vector. The inserted sequence contains the entire coding region of cmEPO (Genbank accession M18189). See SEQ ID NO:3.

#### B. Mice

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Immunodeficient severe combined immunodeficiency (SCID) mice were obtained from Charles River Labs, Wilmington, Massachusetts, USA.

Intramuscular injections were performed as follows: mice were anaesthetized with 50 µl of a solution which contained 80 mg/ml ketamine and 4 mg/ml of xylazine. The area surrounding the anterior tibialis muscle was shaved. Fifty µl of DNA, at a concentration of 2.7 ug/µl in .9% saline solution was injected into the anterior tibialis muscle of both legs using a 28 gauge needle. Twenty-four hours after the first injection, a second injection was performed using the identical protocol. Blood was taken from the orbital sinus to determine hematocrits on a weekly basis.

#### C. Result

The hematocrit readings on 6 mice which were injected with plasmid, CMVkm-cmEpo (which expresses the cynomolgus monkeys EPO cDNA), are shown in Table 3 below. The row marked control shows the average reading for three uninjected mice. The raw data for the three control mice is shown in the lower part of Table 3. Mouse 2 in the injected group died between 4 and 5 weeks post-injection.

Table 3

,						Hemato	Hematocrit Levels (%)	<b>[9</b> ]				
	Week#	Week 0	week I	week 2	week 3	week 4	week 5	week 6	week 7	week8	week 9	week 10
	ſì									Marine Section 201		
	Mouse ⊍											
	mouse 1	50	63	66.5	57.5	63	63.5	56.5	62.5	54	53.5	545
	mouse 2	50	64	49	56.5	55.5						
	mouse 3	50	09	61.5	63	61	56	49.5	53	53.5	54.5	57
<u>-</u>	mouse 4	50	62	68.5	71.5	67.5	99	62.5	59.5	57.5	53.5	555
	mouse 5	50	62	62.5	56	19	53.5	58	52.5	54	5 65	48.5
60	mouse 6	50	99	63.5	62.5	09	58	58	53.5	55.5	\$ 65	505
7	Control	50	51.5	48	47.5	53	49.5	49.5	50.5	515	51	49
l.												
******												
	control	week 0	control	control	control	control	control	control	control	control	control	control
ىلىب			wk 1	wk 2	wk3	week 4	week 5	week 6	week7	week 8	week 9	week 10
	mouse I	20	52	48	46.5	52.5	48	49.5	52	51.5	505	46.5
		50	51	47	47	55	51	50.5	50	52	49	515
	mouse 3	50	52	49	48.5	52	49.5	48.5	48 5	21	505	50

#### Example 8

#### Injection of Polynucleotides Encoding Leptin

#### A. Polynucleotides

The CMV-km2 vector, described above, was used for these experiments. Either the wild-type or HA version of the leptin coding region was inserted into the vector. The map of the plasmid is depicted in Figure 4 and the sequence of the vector with the wild type leptin is shown in SEQ ID NO:4.

#### B. Mice

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Ob/ob mice were obtained from Jackson Labs, Bar Harbor, Maine, USA. The first of the recessive obesity mutations, the *obese* mutation (*ob*) was identified and described in 1950 by Ingall et al., 1950, J. Hered. 41:317-318. Subsequently, 5 single-gene mutations in mice have been observed to produce an obese phenotype, as described in Friedman et al., 1990, Cell 69:217-220. (More recently, the mouse *obese* gene and its human homologue have been cloned, as described in Zhang et al., 1994, Nature 372:425).

#### C. Method

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Intramuscular injections were performed as follows: mice were anaesthetized with the same ketamine solution described above in the Example 7 and the area surrounding the anterior tibialis muscle was shaved.

Fifty microliters of DNA at a concentration of 3.3.  $\mu$ g/ $\mu$ l in 0.9% saline solution was injected into the anterior tibialis muscle of both legs using a 28 gauge needle.

Seventy-two hours after the first injection, a second injection was performed using the identical protocol.

Group 1 ob/ob mice were injected with a plasmid (CMVkM leptin-wt) which encodes the wild-type mouse leptin protein.

Group 2 ob/ob mice were injected with a plasmid (CMVkm-leptinHA) which encodes a form of mouse leptin which is modified with the epitope which is recognized by the antibody 12CA5. The amino acid sequence of the epitope is SYPYDVPDYASLGGPS (Wilson et al., 1984, Cell 37: 767-778).

Group 3 ob/ob mice were injected with a solution of 0.9% saline.

The mice were weighed each day (see Table 4) and the proportional weight gain for each mouse during the first eight days was calculated. The results are shown in Table 5. For any given day, the weight was subtracted from the weight of the individual mouse on day 0, and the difference was divided by the weight on day 0. The proportional weight change data from day 8 was analyzed using an unpaired t-test. When compared with group 3 control mice the p value from group 2 mice was 0.004. When compared with group 3 control mice, the p value for group 1 mice is 0.0038.

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Note: the mice were not weighed on day 1 and day 2, the values for these days were extrapolated from day 3.

Table 4

Weight	Weight of Mice in Grams	n Gram	S								
group 1	day 0	day3	day4	day5	dav6	dav7	day.e	9000	3.		
mouse 1	47	40	ę	5	1				dayio	day11	day 12
	┿			5			?	25	52	53	53
7 asmorri	╅	44	49	51	51	51	51	52	53	\$	55
mouse 3	46	48	48	49	49	49	40	Ş	3 3	75	53
mouse 4	47	48	37	ę	١			2	2	20	51
3 001100	= =		٤		÷	4	8	20	20	50	51
Casnolli	44	2	2	2	51	51	52	52	52	52	3
											*
group 2	day 0	day 3	dav 4	day 5	day 6	1000					
mouse	40	5				, leave	o kgo	day y	day 10		
		3	3	75	77	52	52	53	54	7.5	55
mouse I	43	45	4	45	45	4	45	45	¥	5	3 1
mouse 3	48	49	49	ę,	ç	S	2 3	7 8	⊋ ;	40	47
monse 4	9	Ş	5		3	3	7	77	52	52	52
. 2000		3	2	77	2	51	<b>2</b> 5	52	52	53	53
mouse 5	<del>2</del>	48	49	49	49	50	50	51	15	* 5	3
		-							1	7	21
group 3	day 0	dav 3	day 4	day 5	400.6						
monse 1	8					(ay	day 8	day 9	day 10		
	2 5	2	77	£	43	45	45	45	45	45	46
mouse 2	48	49	50	50	51	52	53	53	:   5	$\dagger$	2
mouse 3	48	50	50	52	53	53	:   2	†	77		53
monse 4	40	53	5	1 2	3 8		3	2	55	55	55
3 00000	1	7	† */:	3	2	74	55	25	54	5.4	55
Casponi	5	\$	8	47	48	46	50	49		T	3 5
									_		7

Table 5

Proportio	anal Cha			D					
Proportion (gp1), Le	mai Chai	ige in w	eignt ire	m Day u	of Mice	Injected	with cE	NA for	Leptin
(gp1), Le	pun-ma	(gpz) or	Same						
Group I	Mice Be	low Injec	ted with	CMVIA	A-I antir				··········
	day 0	day I	day 2	day 3	day 4	day 5	1	T	T
mouse 1	0	0.009	0.018	0.028	0.028		day 6	day 7	day 8
mouse 2	0	0.003	0.026	0.028	<del></del>	0.059	0.059	0.059	0.066
mouse 3	0	0.006	0.020	0.039	0.039	0.085	0.085	0.085	0.085
mouse 4	0	<del></del>	<del></del>	<del></del>	0.018	0.062	0.062	0.062	0.062
mouse 5	0	0.014	0.028	0.042	0.042	0.065	0.065	0.065	0.065
mouse 3	0	0.007	0.014	0.021	0.021	0.042	0.042	0.042	0.06
	10-	0.007	0.014	0.021	0.021	0.041	0.041	0.041	0.061
Group 2.1	Mice Rel	L Inica	المنامل	CMAN	<u> </u>	<u> </u>	<u> </u>		<u> </u>
Group 2 1								·	<del></del>
mouss I	day 0	day 1	day 2	day 3	day 4	day 5	day 6	day 7	day 8
mouse i	0	0.006	0.013	0.02	0.02	0.06	0.06	0.06	0.06
mouse 2	0	0.013	0.031	0.046	0.023	0.046	0.046	0.046	0.046
mouse 3	<u> </u>	0.007	0.014	0.021	0.021	0.04	0.04	0.04	0.062
mouse 4	0	0.006	0.012	0.02	0.02	0.06	0.04	0.04	0.06
mouse 5	0	0.014	0.028	0.043	0.065	0.065	0.065	0.08	0.08
		L		<u> </u>	<u> </u>	<u> </u>			
Group 3 1						<del></del>	·····		
	day 0	day 1	day 2	day 3	day 4	day 5	day 6	day 7	day 8
mouse I	0	0.016	0.032	0.05	0.05	0.06	0.06	0.125	0.125
mouse 2	0	0.07	0.014	0.021	0.04	0.04	0.06	0.08	0.08
mouse 3	0	0.013	0.026	0.04	0.04	0.08	0.08	0.1	0.1
mouse 4	0	0.02	0.04	0.06	0.06	0.08	0.08	0.1	0.12
mouse 5	0	0.015	0.03	0.045	0.069	0.09	0.12	0.14	0.14
Average I				ight for	Each Gr	oup			
	day 0	day 1	day 2	day 3	day 4	day 5	day 6	day 7	day 8
group 1	0	0.009	0.018	0.028	0.028	0.059	0.059	0.059	0.066
group 2	0	0.009	0.018	0.028	0.029	0.054	0.05	0.053	0.061
group 3	0	0.014	0.028	0.042	0.051	0.07	0.08	0.109	0.113

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#### Example 9

# Peptoid Mediated In Vitro Delivery in COS, HT1080, and 293 Cell Lines

COS cells (available from the American Type Culture Collection, Rockville, MD, under Accession No. CRL 1651 and HT1080 cells (available from the American Type Culture Collection, Rockville, MD, under Accession No. CCL 121) were cultured and transfected with pCMVkmLUC and various polycationic agents of the present invention (described in Example 1) according to the transfection protocol described in Example 4. Luciferase activity was assayed according to the method described in Example 4. Total cell protein was measured using a Pierce BCA kit according to manufacturer's directions.

The results, shown in Figure 7A, indicate that the ability of the polycationic agents to mediate transfection is not dependent on cell line type. Polycationic agents having a repeating trimer motif of neutral and cationic sidechains were particularly effective at mediating transfection.

Transfection efficiencies for a homologous series of cationic peptoids were evaluated. Specifically, cationic peptoids RZ-110-1 (18-mer), RZ-120-3 (24mer), RZ120-7 (36mer), and RZ120-11 (48mer), which have the same repeating (HHP) motif were evaluated for their ability to transfect COS and HT1080 cells. These polycationic agents were complexed with pCMVkmLUC at a 2:1, + to - charge ratio. The concentration of negative charges on DNA was calculated using 3.03 nmol of phosphate per 1 µg of DNA, on the basis of the average molecular weight of 330 for each nucleotide. The formula weight of the polycationic agent was calculated as a semi-trifluoroacetate salt (50% of amino groups form salt with TFA), and the concentration of the polycationic agent was determined on the basis of the weight of the lyophilized peptoid. Amino groups were formally considered to be fully protonated to obtain the number of positive charges on the polycationic agent interest when calculating the + to - charge.

As shown in Figure 7B, transfection efficiencies for this particular series of cationic peptoids were largely independent of oligomer length for peptoids having 24 or more monomeric units.

Transfection efficiencies using polycationic agent RZ145-1 and commercially available cationic lipids, DMRIE-C[™], Lipofectin[®] and lipofectamine were evaluated. In these experiments RZ145-1 was complexed with pCMVkmLUC at a 2:1, + to - charge ratio.

Transfection with DMRIE-C[™] Lipofectin[®], lipofectamine was conducted according to

manufacturer's directions. The cationic lipids were also employed at a 2:1, + to - charge ratio. 293 human embryonic kidney cells (Microbix, Toronto, Ontario, Canada), HT1080 cells, and NIH-3T3 cells (available from the American Type Culture Collection, Rockville, MD, Accession No. CRL 1658) were transfected, cultured either in the presence or absence of 10% serum, then assayed for luciferase production using the same protocol as described in Example 4. Luciferase was measured, as described in Example 4, 48 hours after initial transfection. Total cell protein was measured using a Pierce BCA kit according to manufacturer's directions.

The results, shown in Figure 8, indicate that, in contrast to Lipofectin® and lipofectamine, which were respectively 10- and 100-fold less efficient in the presence of serum, gene transfer mediated by polycationic agent RZ145-1 was insensitive to the presence of serum.

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Transfection mediated by polycationic polymers, such as polylysine and histones, is greatly enhanced by addition of chloroquine to the transfection media. To determine whether chloroquine affected transfection mediated by polycationic agents of the present invention, HT1080 and 293 cells were transfected using RZ145-1 in the presence and absence of chloroquine. As a control, the same cell lines were transfected with polylysine both in the presence and absence of chloroquine. The results, shown in Figure 9, indicate that the polycationic agent RZ145-1 was equally effective at mediating transfection both with and without chloroquine. In contrast, polylysine-mediated transfection in the absence of chloroquine was 100-fold lower than polylysine mediated transfection in the presence of chloroquine. In addition, the results indicate that cationic peptoid mediated transfection is more efficient than polylysine mediated transfection.

#### Example 10

Preparation of a Stable Formulation of DNA/Polycationic Agent Complex

A. DNA/Polycatinic Agent Complex Formation (2:1, + to - Change Ratio)

All operations were carried out at ambient temperature. DGPW (diagnosis grade purified water) was used to prepare the stock solutions. Both the plycatinic agent and DNA samples had low salt concentrations (i.e., < lmM) to avoid precipitation.

### (1) Batch Method

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Complexes of polycationic agent RZ145-1 and pCMVkmLUC, as follows, for up to 250 µg DNA. DNA (i.e., pCMVkmLUC) was diluted with 30% (v/v) ethanol in water to a concentration of 50 µg/ml corresponding to 151 µM of negative charge. RZ145-1 was diluted to 23.2 µM in 30% ethanol in water. To 1 part of the polycationic agent solution was added 1 part of DNA solution as quickly as possible with gentle agitation. The DNA solution was added to the solution of polycationic agent (rather than vice-versa) to avoid precipitation. Slow addition of the two solutions was avoided to avoid precipitation and the formation of large complexes.

#### (2) Continuous Method

For more than 250 µg of DNA, a continuous method for preparing concentrated formulations of polycationic agent/DNA complex is preferred. The DNA and peptoid solutions were prepared as above and placed into separate bottles. Each bottle was connected to one port of a mixing tee. The bottles were simultaneously pressurized with 2 to 3 psi to deliver the two streams to the mixing tee at the same flow rate (e.g., 20 ml/min or higher).

### B. Concentration Step

Two milliliters of the DNA-polycationic agent complex from part A was placed in a Centricon*-100 (Amico Inc. Beverly, MA), and centrifuged at 1000 x g for 30 minutes or until the volume of the retentate containing polycationic agentDNA complex was approximately 50 µl. The filtrate was removed from the bottom receiver. The retentate was diluted with 2 ml of 5% glucose, and concentrated to 50 µl again. This operation was repeated again to produce a concentrated complex solution containing 1 mg/ml DNA in 5% glucose. This concentration step can be conducted at either 4°C or at ambient temperature. The ethanol content of the final concentrated solution was less than 0.1%. No precipitation was observed in the concentrated solution.

#### SEQUENCE LISTING

- (1) GENERAL INFORMATION:
  - (i) APPLICANT: Zuckermann et al.
- (ii) TITLE OF INVENTION: Compositions and Methods for Polynucleotide Delivery
- (iii) NUMBER OF SEQUENCES: 4
- (iv) CORRESPONDENCE ADDRESS:
  - (A) ADDRESSEE: Chiron Corporation
  - (B) STREET: 4560 Horton Street
  - (C) CITY: Emeryville
  - (D) STATE: California
  - (E) COUNTRY: U.S.A.
  - (F) ZIP: 94608-2916
  - (v) COMPUTER READABLE FORM:
    - (A) MEDIUM TYPE: Floppy disk
    - (B) COMPUTER: IBM PC compatible
    - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
    - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
  - (vi) CURRENT APPLICATION DATA:
    - (A) APPLICATION NUMBER:
    - (B) FILING DATE:
    - (C) CLASSIFICATION:
  - (viii) ATTORNEY/AGENT INFORMATION:
    - (A) NAME: Fujita, Sharon M.
    - (B) REGISTRATION NUMBER: 38,459
    - (C) REFERENCE/DOCKET NUMBER: 1218.002
    - (ix) TELECOMMUNICATION INFORMATION:
      - (A) TELEPHONE: (510) 923-2706
      - (B) TELEFAX: (510) 655-3542
- (2) INFORMATION FOR SEQ ID NO:1:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 9600 base pairs
    - (B) TYPE: nucleic acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: DNA (genomic)
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

GCGGCCGCGG	AATTCTCATG	TTTGACAGCT	TATCATCGAT	AAGCTGATCC	TCACAGGCCG	60
CACCCAGCTT	TTCTTCCGTT	GCCCCAGTAG	САТСТСТСТС	TECTEACCTT	GAAGAGGAAG	• •
AGGAGGGGTC	CCGAGAATCC	CCATCCCTAC	CCMCCTCTCTC	IGGIGACCII	GAGGAATTTG	120
ACCCCTCCCC	TO COMMENCE	CONTCCCIAC	CGTCCAGCAA	AAAGGGGGAC	GAGGAATTTG	180
MOGCCIGGCI	TGAGGCTCAG	GACGCAAATC	TTGAGGATGT	TCAGCGGGAG	TTTTCCGGGC	240
TGCGAGTAAT	TGGTGATGAG	GACGAGGATG	GTTCGGAGGA	TGGGGAATTT	TCAGACCTGG	300
ATCTGTCTGA	CAGCGACCAT	GAAGGGGATG	AGGGTGGGGG	GGCTGTTGGA	GGGGCAGGA	
GTCTGCACTC	CCTGTATTCA	CTGAGCGTCG	TCTAAMAAAC	MCCCCTTTTT	ATCTCTTTTA	360
GTGTGAATCA	TOTOTO A COL	000000000000000000000000000000000000000	TOTANTANA	ATGTCTATTG	ATCTCTTTTA	420
CCACAGAGAG	IGICIGACGA	GGGGCCAGGT	ACAGGACCTG	GAAATGGCCT	AGGAGAGAAG	480
GGAGACACAT	CTGGACCAGA	AGGCTCCGGC	GGCAGTGGAC	CTCAAAGAAG	AGGGGGTGAT	540
AACCATGGAC	GAGGACGGG	AAGAGGACGA	GGACGAGGAG	GCGGAAGACC	AGGAGCCCCC	600
GGCGGCTCAG	GATCAGGGCC	AAGACATAGA	GATICATION	CCACACCCCA	11110000000	•
AGTTGCATTG	GCTGCAAAGG	CACCCACCOM	COLLEGIGICC	GGAGACCCCA	AAAAUGTCCA	660
	GCIGCAAAGG	GACCCACGGT	GGAACAGGAG	CAGGAGCAGG	AGCGGGAGGG	720

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GCAGGAGCAG	CACCCCCAC					
GCAGGAGGGG	CAGGGGGCAGG	AGCAGGAGGA	GGGGCAGGAC	G CAGGAGGAGG	G GGCAGGAGGG	780
GGGGCAGGAG	GCCACCACC	AGGAGCAGGA	GGAGGGGCAG	GAGCAGGAGG	G GGCAGGAGGG G AGGGGCAGGA	840
GGAGCAGGAG	CACCCCCA	AGGAGGAGGG	GCAGGAGCAC	GAGGAGGGG	AGGGGCAGGA  AGGAGGGGCA	900
GGAGGAGGAG	GAGGGGCAGG	AGGGGCAGGA	GGGGCAGGAC	G CAGGAGGAGG	GGCAGGAGCA	960
DODDADDADD.	CAGGAGGGGC	AGGAGCAGGA	GGAGGGGCAG	GAGGGGCAG	GGCAGGAGCA  GAGGGGCAGGA	1020
GCNGGNGGNG	GGGCAGGAGC	AGGAGGGCA	GGAGGGGCAG	GAGGGGGCACC	1 1001001000	1080
OCHOOMOCAG	GAGGAGGGGC	AGGAGGGCA	GGAGGGGCAC	GAGCAGCACC		1140
OCHOOGICAG	GAGCAGGAGG	GGCAGGAGCA	GGAGGGGCAG	GAGGGGGCACC	100100100	1200
CHOCHOOGG	CAGGAGCAGG	AGGGGCAGGA	GGGGCAGGAG	CAGGAGGAGG	0003003000	
OKJORODROCAG	GAGGAGGGG	AGGAGGGCA	GGAGCAGGAG	GGGCAGGAGG		1260
CONGCOCAG	GAGGGGCAGG	AGCAGGAGGG	GCAGGAGGGG	CAGGAGGAGG	10010000	1320
ONOCAGOAG	GGGCAGGAGC	AGGAGGTGGA	GGCCGGGGGTC	GAGGAGGGAG	moot coopee	1380
GGTCGAGGAG	GTAGTGGAGG	CCGGGGTCGA	GGAGGTAGTG	GAGGGCCCC	TGGAGGCCGG	1440
CGTGAAAGAG	CCAGGGGGG	AAGTCGTGAA	AGAGCCACCC	CANGGLEGEEG	GGGTAGAGGA	1500
GAAAAGAGGC	CCAGGAGTCC	CAGTAGTCAG	TCATCATCAT	GGAGAGGTCG	TGGACGTGGA	1560
CCCCCTCCAG	GTAGAAGGCC	Puninimuccyc	CCMCCALCAL	CCGGGTCTCC	ACCGCGCAGG	1620
CACCAAGAAG	GTGGCCCAGA	TEGTENCE	CACCTGTAGGGG	AAGCCGATTA	TTTTGAATAC	1680
CCCGCAGATG	ACCCAGGAGA	ACCCCCAACC	ACCORDING	CGGGAGCGAT	AGAGCAGGGC	1740
AGGCGCAAAA	ACCCAGGAGA AAGGAGGGTG	GOODCOAAGC	ACTGGACCCC	GGGGTCAGGG	TGATGGAGGC	1800
TTTGAGAACA	AAGGAGGGTG	TTTT ACT COM	CATCGTGGTC	AAGGAGGTTC	CAACCCGAAA	1860
ACCGACGAAG	TTGCAGAAGG	CCCCCCCCCC	CTCCTGGCTA	GGAGTCACGT	AGAAAGGACT	1920
TACAACCTAA	GAACTTGGGT	CGCCGGTGTG	TTCGTATATG	GAGGTAGTAA	GACCTCCCTT	1980
CCTCTCCCCC	GGCGAGGAAC	TGCCCTTGCT	ATTCCACAAT	GTCGTCTTAC	ACCATTGAGT	2040
ATTOCCCCT	TTGGAATGGC	CCCTGGACCC	GGCCCACAAC	CTGGCCCGCT	AAGGGAGTCC	2100
*** ******	ATTICATEGI	CTTTTTACAA	ጥልጥልጥልግግል	THE COMPANIES	mmma	2160
occur I wwo	ACCTTGTTAT	GACAAAGCCC	COMPONDATION	CCS AMB MOS O	00mos	2220
***************************************	ACGATGGAGT.	AGATTTGCCT	CCCTCCTTTC	CACCMAMOOM	CO11 CO000	2280
Controcation	GIGMIGACCG	AGATGACGGA	GATGAAGGAG	CTCATCATCA	Maranes	2340
O. B. GOOCAGO	AGIGATGTAA	CTTGTTAGGA	GACGCCCTCA	ATCOMAMMAA	******	2400
TTCCCCCGCM	CIMMAGAATA	AATCCCCAGT	AGACATCATG	CGTGCTGTTC	CECENTER	2460
COCCATCIGI	CITGICACCA	TTTTCGTCCT	CCCAACATGG	CCCBATTTCCC	CRMBCCCTBC	2520
TIGICACGIC	ACTUAGCTCC	GCGCTCAACA	CCTTCTCGCG	TTCCAAAAAA	MM1 0001 01 0	2580
* 1 MCC 1 GG 1 G	AGCAATCAGA	CATGCGACGG	CTTTAGCCTG	GCCTCCTTA A	A MMC A COMA A	2640
OUNT GOOD WOC	MACCAGCATG	CAGGAAAAGG	ACAAGCAGCG	BABAMMORGO	~~~~~~~	
NGG 1 GG C GG C	ATATGCAAAG	GATAGCACTC	CCACTCTACT	A CTCCCCTTA TO	1 m1 maama1 -	2700
TOTATATGCA	TGAGGATAGC	ATATGCTACC	CGGATACAGA	TTACCAMACO	3 77 3 77 3 77 7 7	2760
CHONTHINGH	TTAGGATAGC	ATATGCTACC	CAGATATAGA	TTACCAMAGG	C/13 mccms s-	2820
CHONINIMAN	TTAGGATAGC	ATATACTACC	CAGATATAGA	TTACCAMACC	3.003.000×00	2880
CHONININGN	TTAGGATAGC	CTATGCTACC	CAGATATAGA	TTAGGAMAGG	3.003.000 a.c.	2940
CUCKININGN	TTAGGATAGC	ATATGCTATC	CAGAMAMMMC	CCTACTACA	GGM3 GGM4	3000
TATAAATTAG	GATAGCATAT	ACTACCCTAA	ТСТСТАТТАС	CATACCAMAM	GCTACCCAGA	3060
TACAGATTAG	GATAGCATAT	ACTACCCAGA	ጥልጥልርልጥጥልር	CVEVCCVIVI	GCTACCCGGA	3120
TATAGATTAG	GATAGCCTAT	GCTACCCAGA	TATALATTAC	CATAGCATAT	GCTACCCAGA	3180
TATAGATTAG	GATAGCATAT	GCTACCCAGA	TATACATTAC	CATAGCATAT	ACTACCCAGA	3240
TATAGATTAG	GATAGCATAT	GCTATCCAGA	TATAGATIAG	GATAGCCTAT	GCTACCCAGA	3 <b>30</b> 0
CATTAGCCCA	CCGTGCTCTC	ACCGACCTCC	MCI INCIDENT	GTATATGCTA	CCCATGGCAA	3360
GCGCTCAGGC	GCAAGTGTGT	CONTRACTOR	TGAATATGAG	GACCAACAAC	CCTGTGCTTG	3420
TGGCCCGCCC	ACCTACTTAT	CCYCCAYAAA	CTCCAGATCG	CAGCAATCGC	GCCCCTATCT	3480
AGTGGTTTGA	ACCTACTTAT	TACCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	CCCGGGGTGC	CATTAGTGGT	TTTGTGGGCA	3540
TACAGTCCAA	CCGCAGTGGT	TAGCGGGGTT	ACAATCAGCC	AAGTTATTAC	ACCCTTATTT	3600
CAAAAAAAA	AACCGCAGGG	CGGCG1G1GG	GGGCTGACGC	GTGCCCCCAC	TCCACAATTT	3660
AATTTTCGGG	AGTGGCCACT	AGAAGAAGA	TATGGGCCCC	ATTGGCGTGG	AGCCCCGTTT	3720
TCCCCTTCTT	GGTGTTAGAG	ACAACCAGTG	GAGTCCGCTG	CTGTCGGCGT	CCACTCTCTT	3 <b>78</b> 0
CACATCTTA	ACAAATAGAG	TGTAACAACA	TGGTTCACCT	GTCTTGGTCC	CTGCCTGGGA	3840
OUCULC LIVY	TAACCCCAGT	ATCATATTGC .	ACTAGGATTA	TGTGTTGCCC	ATAGCCATAA	3900
Witcold GM	GATGGACATC	CAGICITIAC -	GGCTTGTCCC	CACCCCAMCC	a mmmoma mmo	3960
ITAAAGATAT	TCAGAATGTT '	TCATTCCTAC .	ACTAGTATTT	<b>ል</b> ጥጥርርርርር እአር	CCCMMmamama	4020
GGGIIAIAI.L	GGTGTCATAG	CACAATGCCA (	CCACTGAACC	CCCCCCCCC	1 mmmn	4080
GGGGGCGTCA	CCIGAAACCT '	TGTTTTCGAG	ርልሮሮጥሮኔሮኔሞ	ACACCMMACM	~~~~~~~~~	4140
CAGCAGIIAT	TCTATTAGCT	AAACGAAGGA	GAATGAAGAA	GCAGGCCAAC	3 MMC 3 CC 2 C 2	4200
GII CVC I GCC	CGCTCCTTGA '	TCTTCAGCCA (	CTGCCCTTGT	GACTAABATC	COMPORTOR OF	4260
CICOIGGNAI.	CCTGACCCCA '	TGTAAATAAA	ACCGTGACAG	しぶし かむいへいしゅ	0001010100	4320
derettect.L.	AGGACCCTTT (	TACTAACCCT .	AATTCGATAG	こののことである。	COOMMOOGEN	4320
HCMINIGCIA.	TIGAATTAGG (	GTTAGTCTGG .	ATAGTATATA	CTACTACCC	~~**	
GCTACCCGTT	TAGGGTTAAC	AAGGGGGCCT '	TATAAACACT	ATTGCTAATC	CCCMCMMC*C	4440
			78	nn10	CCCICIIGAG	4500

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GGTCCGCTT	A TCGGTAGCT	A CACAGGCCC	C TCTGATTGA	C GTTGGTGTA	G CCTCCCGTAG	4560
TCTTCCTGG	G CCCCTGGGA	G GTACATGTC	C CCCAGCATT	G GTGTAAGAG	C TTCAGCCAAG	4620
AGTTACACA	T AAAGGCAAT	G TTGTGTTGC	A GTCCACAGC	A TGCAAAGTC	T GCTCCAGGAT	4680
CACAAGCCAC	T CAGTGTTGG	CAAATGTGCA	C ATCCATTTA	P AAGGATGTC.	A ACTACAGTCA	4740
AGTTGTACC	Trigrary rg	TCCCCCCCC	G TGTCACATG	r ggaacaggg	C CCAGTTGGCA	4800
CCTCGTACC	A CCCAACTGA	A GGGATTACA	r GCACTGCCC	GAATACAAA	A CAAAAGCGCT	4860
GGGCGATCG	R GCGAAGAAGG	GGCAGAGAT(	G CCGTAGTCAG	G GTTTAGTTC	G TCCGGCGCG	4920
GGCGTCGGC	CACCOUCT(	CGCGCTCGC	r CGCTCACTG	A GGCCGCCCG	G GCAAAGCCCG	4980
TGGCCAACT	CARCUITIGG	r CGCCCGGCC	r CAGTGAGCG	A GCGAGCGCG	C AGAGAGGGAG	5040
CTACGGCCC	CATCACTAG	GGTTCCTTGT	r AGTTAATGAT	TAACCCGCC	A TGCTACTTAT	5100
TACATTTACT	CGARTITUG	A CTCTAGGCCA	A TTGCATACG	r TGTATCTATI	A TCATAATATG	5160
TATTAATAG	T ARTCARCALCALC	CCCCMCAMO	CUGCCATGT	GACATTGAT	T ATTGACTAGT	5220
ACATAACTT	A COGTANATO	GGGGTCATTA	GTTCATAGCO	CATATATGG	ATTORCTAGT	5280
TCAATAATG	CGTATGTTC	COUGCUIGG	. TGACCGCCC	ACGACCCCC	G CCCATTGACG G ACGTCAATGG	5340
GTGGAGTATT	TACGGTAAAC	TOTAL TANGENTAL C	CCAATAGGG	CTTTCCATTC	ACGTCAATGG ATATGCCAAGT	5400
CCGCCCCCT	TTGACGTCAX	TOBCCACILO	TOCCCCCCCCCC	AAGTGTATC	TATGCCAAGT CCAGTACATG	5460
ACCTTACGGC	ACTTTCCTAC	TORCOGIANA	TGGCCCGCCT	GGCATTATGC	CCAGTACATG TATTACCATG	5520
GTGATGCGGT	TTTGGCAGTA	CACCAAMCCC	COMMON	TAGTCATCGC	TATTACCATG ACGGGGATTT	5580
CCAAGTCTCC	ACCCCATTCA	COTTCNATGGG	CGTGGATAGC	GGTTTGACTC	ACGGGGATTT TCAACGGGAC	5640
TTTCCAAAAT	GTCGTAATAA	CCCCCCCCCCC	AGTTTGTTT	GGCACCAAAA	TCAACGGGAC GCGTGTACGG	5700
TGGGAGGTCT	' ATATAAGCAG	AGCTCGTTTA	GTCAACGCAAA	TGGGCGGTAG	GCGTGTACGG GAGACGCCAT	5760
CCACGCTGTT	TTGACCTCCA	TAGAAGACAC	GIGAACCGTC	AGATCGCCTG	GAGACGCCAT CGGCCGGGAA	5820
CGGTGCATTG	GAACGCGGAT	TCCCCGTGCC	' AAGAGTGAGG	CCAGCCTCCC	CGGCCGGGAA CTATAGACTC	5880
TATAGGCACA	CCCCTTTGGC	TCTTATGCAT	GCTATACTCT	TAAGTACCGC	GGGCCTATAC	5940
ACCCCCGCTC	CTTATGCTAT	AGGTGATGGT	ATAGCTTAGC	CTATACCTIC	GGGCCTATAC GGGTTATTGA	6000
CCATTATTGA	CCACTCCCCT	ATTGGTGACG	ATACTTTCCA	TTALTAGGIGI	ATAACATGGC	6060
TCTTTGCCAC	AACTATCTCT	ATTGGCTATA	TGCCAATACT	CTGTCCTTCA	GAGACTGACA	6120
CGGACTCTGT	ATTTTTACAG	GATGGGGTCC	ATTTATTATT	TACAAATTCA	CATATACAAC	6180
AACGCCGTCC	CCCGTGCCCG	CAGTTTTTAT	TAAACATAGC	GTGGGATCTC	CGACATCTCG	6240
GGTACGTGTT	CCGGACATGG	GCTCTTCTCC	GGTAGCGGCG	GAGCTTCCAC	ATCCGAGCCC	6300
TGGTCCCATC	CGTCCAGCGG	CTCATGGTCG	CTCGGCAGCT	CCTTCCTCCT	AACAGTGGAG	6360
GCCAGACTTA	GGCACAGCAC	AATGCCCACC	ACCACCAGTG	TGCCGCACAA	CCCCCMCCCC	6420
GINGGGIAIG	TGTCTGAAAA	TGAGCTCGGA	GATTGGGCTC	GCACCTGGAC	CCACAMCCA	6480
GACTTAAGGC	AGCGGCAGAA	GAAGATGCAG	GCAGCTGAGT	ጥርጥጥርጥልጥጥ	mcamaa.ca.cm	6540
CAGAGGTAAC	TCCCGTTGCG	GTGCTGTTAA	CGGTGGAGGG	CAGTGTAGTC	TO A CO A COM A CO	6600
ICG11GC1GC	CGCGCGCGCC	ACCAGACATA	ATAGCTGACA	GACTAACACA	CITCHING COMMIN	6660
CCMIGGGICI	TTTCTGCAGT	CACCGTCGTC	GACCTAAGAA	TTCAGACTCG	ACCA ACMOMA	6720 6780
GAMAGCCATG	GATATCGGAT	CCACTACGCG	TTAGAGCTCG	CTGATCAGCC	TO A CHICAGO	6840
CITCIAGITG	CCAGCCATCT	GTTGTTTGCC	CCTCCCCCGT	CCCTTCCTTC	ACCOMOGNA	6900
GIGCCACICC	CACTGTCCTT	TCCTAATAAA	ATGAGGAAAT	TOCATOGOAT	mememos ams	6960
GGIGICATTC	TATTCTGGGG	GGTGGGGTGG	GGCAGGACAG	CAAGGGGGAC	CAMMOOOLL	7020
ACAMTAGCAG	GGGGGTGGGC	GAAGAACTCC	AGCATGAGAT	CCCCCCCCCC	CICCIMONA	7080
CAGCCGGCGI	CCCGGAAAAC	GATTCCGAAG	CCCAACCTTT	CATAGAAGGC	CCCCCCCCC	7140
I COMMITTIC	GIGATGGCAG	GTTGGGCGTC	GCTTGGTCGG	ጥር አመመጥር ርርጥ	3 C 3 M 3 3 C M 5 C	7200
CWIGGCGGGT.	TAATCATTAA	CTACAAGGAA	CCCCTAGTGA	TGGAGTTGGC	CACTCACTCA	7260
CIGCOCOCIC	GCTCGCTCAC	TGAGGCCGGG	CGACCAAAGG	TCGCCCCXCC	CCCCCCCC	7320
accedaced	CCTCAGTGAG	CGAGCGAGCG	CGCAGCGAAC	CCCAGAGROO	CCCMCSCSSS	7380
WWC I COT CWW	GAAGGCGATA	GAAGGCGATG	CGCTGCGAAT	CGGGAGGGGG	CAMBAGOMA	7440
AGUACGAGGA	AGCGGTCAGC	CCATTCGCCG	CCAAGCTCTT	CAGCAAMAMO	3.0000000000	7500
MUCGCINIGI	CCTGATAGCG	GTCCGCCACA	CCCAGCCGGC	CACACTCCAM	C11maa.a.	7560
TWGCGGCCAT.	TTTCCACCAT	GATATTCGGC	AAGCAGGCAT	CCCCxmcccm	G10010010	7620
TCCTCGCCGT	CGGGCATGCG	CGCCTTGAGC	CTGGCGAACA	GTTCCCCTCC	00000000	7680
TOWIGGICIT	CGTCCAGATC	ATCCTGATCG	ACAAGACCGG	CTTCCTTCCC	1 CM 1 COMOG	7740
COCICONIGC	GATGTTTCGC	TTGGTGGTCG	AATGGGCAGG	TAGCCCCAMC	******	7800
MOCCOCACA	TIGCATCAGC	CATGATGGAT	ACTTTCTCGG	CAGGAGGAAG	CMC 1 C1 mc1 c	7860
"CONGNICCI	GCCCCGCAC	TICGCCCAAT	AGCAGCCAGT	CCCTTCCCCC	MMC & CMC & C *	7920
ACGI CGAGCA	CAGCIGCGCA	AGGAACGCCC	GTCGTGGCCA	GCCACCAMAC	0000000	7980
* COLCCIOCH	GITCATTCAG	GGCACCGGAC	AGGTCGGTCT	TCACAAAAA	******	8040
2001000010	ACAGCCGGAA	CAUGGCGCA	TCAGAGCAGC	CC 3 TTCTCTCC	mmomos	8100
- CHINGCOM	MARGCCACAC	CACCCAAGCG	GCCGG ACA AC	CTICCOMOGA		8160
	QVVVCQVICC.	TUATCUTGTC	TCTTGATCAG	ልጥሮጥምር አመርር	00000000	8220
	GCGGCAAGAA	AGCCATCCAG	TTTACTTTGC	AGGGCTTCCC	AACCTTACCA	8280
			79			• •

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GAGGGCGCCC	CAGCTGGCAA	TTCCGGTTCG	CTTCCTCTC			
TATCGCCATG		GCAAGCTACC			CCAGTCTAGC	8340
GTCCAGATAG					GTTTTCCCTT	8400
TTCTACGTGT	accurative 1			-1000111010	CGGACTGGCT	8460
GCTGTCAATT		TTAGCAGCCC			GCAGCGTGAA	8520
	AAAAAA TAMMA	TTTTTGTTAA	ATCAGCTCAT	TOTO TOTO TOTO		8580
ATCOGCAAAA	TCCCTTATAA	ATCAAAAGAA	TAGCCCGAGA	TACCCOMOSSO	man	8640
OLLIGGRACA	WOWRITCH WOLL	ATTAAAGAAC	GTGGACTCCA	ACCMON NAME		8700
OTC THICHGG	GCGWIGGCGG	ATCAGCTTAT	GCGGTGTGAA	ATACCCCCACA	GATGCCTAAG	
ALIGHMANT NC	COCATCAGGC	GCTCTTCCGC	TTCCTCGCTC	ACTGACTCCC	TGCGCTCGGT	8760
CGTTCGGCTG	CGGCGAGCGG	TATCAGCTCA	CTCAAAGGCG		TATCCACAGA	8820
ATCAGGGGAT		AGAACATGTG			TATCCACAGA	8880
TAAAAAGGCC	GCGTTGCTGG		TACCCTCCCC	CAGCAAAAGG		8940
AAATCGACGC	TCAAGTCAGA	GGTGGCGAAA			AGCATCACAA	9000
TCCCCCTGGA	AGCTCCCTCG			CTATAAAGAT	ACCAGGCGTT	9060
GTCCGCCTTT			TGTTCCGACC	CTGCCGCTTA	CCGGATACCT	9120
CAGTTCGGTG			GCTTTCTCAT	AGCTCACGCT	GTAGGTATCT	9180
CGACCGCTGC	TAGGTCGTTC		GGGCTGTGTG	CACGAACCCC		9240
	GCCTTATCCG		TCTTGAGTCC	AACCCGGTAA	GACACGACOO	9300
ATCGCCACTG	GCAGCAGCCA		GATTAGCAGA	GCGAGGTATG	TAGGCGCTCC	
		GGCCTAACTA	CGGCTACACT	AGAAGGACAG	TATTTCCTA	9360
CTGCGCTCTG		TTACCTTCGG	AAAAAGAGTT	~~~		9420
	GCTGGTAGCG		GTTTGCAACC	AGCAGATTAC	GATCCGGCAA	9480
AAAGGATCTC	AAGAAGATCC	TTTGATCTTT	TCTTACTGAA	CCCTCATTAC	GCGCAGAAA	9540
			I IIIC I GAA	CGGTGATCCC	CACCGGAATT	9600

# (2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 4328 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

GCCGCGGAAT	TTCGACTCTA	GGCCAMMOOA	m>			
TTATATTGGC	TCATGTCCAA	TAMEN GOOG	TACGTTGTAT	CTATATCATA	ATATGTACAT	60
ATAGTAATCA	ATTACGGGGT	CAMMACMMA	ATGTTGACAT	TGATTATTGA	CTAGTTATTA	120
ACTTACGGTA	ATTACCCCCC	CATTAGTTCA	TAGCCCATAT	ATGGAGTTCC	GCGTTACATA	180
AATGACGTAT	GTTCCCATAC	UTGGCTGACC	GCCCAACGAC	CCCCCCCAT	TGACGTCAAT	240
GTATTTACGG	TAAACTCCCATAG	TAACGCCAAT	AGGGACTTTC	CATTGACGTC	AATGGGTGGA	300
CCCTATTGAC	GTCA ATCACC	ACTIGGCAGT	ACATCAAGTG	TATCATATGC	CAAGTCCGCC	360
ACGGGACTTT	CCMACMMOOG	GTAAATGGCC	CGCCTGGCAT	TATGCCCAGT	ACATGACCTT	420
GCGGTTTTCC	CACMACATGGC	AGTACATCTA	CGTATTAGTC	ATCGCTATTA	CCATGGTGAT	480
TCTCCACCCC	AMMORGONO	ATGGGCGTGG	ATAGCGGTTT	GACTCACGGG	GATTTCCAAG	540
DODOCACEC A & & &	ATTGACGTCA	ATGGGAGTTT	GTTTTGGCAC	CAAAATCAAC	GGGACTTTCC	600
immuticate of	MATAACCCCG	CCCCGTTGAC	GCAAATGGGC	CGTACCCCCC	M3.000maa.a.	660
GGICIAIAIA	AGCAGAGCTC	GTTTAGTGAA	CCGTCAGATC	GCCTCCAGAG	000100	720
CIGILLIGAC	CICCATAGAA	GACACCGGGA	CCGATCCAGC	CTCCCCCCCC	00033000	780
Carr + OGMACG	COGMITTECC	GTGCCAAGAG	TGACGTAACT	ACCCCCCCCAA	03 cmc======	840
GCUCUCCCCI	TTGGCTCTTA	TGCATGCTAT	ACTGTTTTTC	COMMOCOCO	Ma Ma ma ma ma	900
COCICCITAL	GCTATAGGTG	ATGGTATAGC	TTAGCCTATA	GGTGTCCCCOO	1 mmc 1 co	960
111 TONCCACI	CCCCTATIGG	TGACGATACT	TTCCATTACT	A A TO CO A TO A CO	1000000	1020
OCCACAMCIA	TUTUTATTGG	CTATATGCCA	ATACTCTGTC	CTTCACACAC	MC 1 C 1 C C	1080
TOTOTALLI	TACAGGATGG	GGTCCATTTA	ፈ ፈ ጉል ምጥጥ ልጥጥ	ልጥጥሮ አር አጠልm	101101100	1140
COLCCCCCC	GCCCGCAGTT	TTTATTAAAC	ATAGCGTGGG	BTCTCCCBCB	mamaaaaa	1200
GIGITCCOOM	CATGGGCTCT	TCTCCGGTAG	CGGCGGAGCT	TOCACATOCC	1.CCCCCCCCC	1260
CCMICCGICC	AGCGGCTCAT	GGTCGCTCGG	CAGCTCCTTC	CTCCTAACAC	MCC1 0000	
MOLLHOGCAC	AGCACAATGC	CCACCACCAC	CAGTGTGCCG	CACAACCCCC	Magaaaaa aa	1320
GINIGICI	GAMMATGAGC	TCGGAGATTG	GGCTCGCACC	TGGACCCACA	mccs s cs com	1380
DODNODIA	CAGAAGAAGA	TGCAGGCAGC	TGAGTTGTTG	ጥ እ መጥረጥረ እ ጠ እ	1010000000	1440
GINNCICCC	TIGCGGTGCT	GTTAACGGTG	GAGGGCAGTG	TACTOTO ACC	1001000	1500
961966666	GCGCCACCAG	ACATAATAGC	TGACAGACTA	ACACACMOMM	acommon a a a a a	1560
0010111101	GCAGTCACCG	TUGTEGACET	AAGAATTCAG	ACTOGRACIONA	CMCMAGAAAA	1620
CCMIGGAINT	CGGATCCACT	ACGCGTTAGA	GCTCGCTGAT	CAGCCTCCAC	mcmcaammam	1680
MOLIGCEMGE	CATCTGTTGT	TTGCCCCTCC	CCCGTGCCTT	CCTTCACCC	001100000	1740
ACTCCCACTG	TCCTTTCCTA	ATAAAATGAG	GAAATTGCAT	CGCATTGTCT	CACHAGGTGCC	1800
			Ø A	occurrency.	GAGTAGGTGT	1860

80

CATTCTATT	C TGGGGGGTGC	G GGTGGGGCAC	GACAGCAAG	G GGGAGGATTO	GGAAGACAAT	1920
AGCAGGGGG	G TGGGCGAAGA	ACTCCAGCAT	GAGATOCCC	G CGCTGGAGG	TCATCCAGCC	1980
GGCGTCCCG	G AAAACGATTO	CGAAGCCCAA	CCTTTCATA	G AAGGCGGCGC	TGGAATCGAA	2040
ATCTCGTGA!	r ggcaggttgc	GCGTCGCTT	GTCGGTCAT'	T TOGAACCCC	CACTCCCCC	2100
CAGAAGAAC:	r cgtcaagaac	GCGATAGAAG	GCGATGCGC	T GCGAATCGGG	ACCCCCCAMA	2160
CCGTAAAGC	A CGAGGAAGCG	GTCAGCCCAT	* TCGCCGCCA	A GCTCTTCAGC	* A A TA TO CA COO	2220
GTAGCCAACC	F CTATGTCCTG	ATAGCGGTCC	GCCACACCC	A GCCGGCCACA	CTCCATCA Am	2280
CCAGAAAAGC	GGCCATTTTC	: CACCATGATA	TTCGGCAAG	AGGCATCGCC	ATCCCTC ACC	2340
ACGAGATCCT	r ceccerces	CATGCGCGCC	TTGAGCCTGG	G CGAACAGTTC	CC TCCCCCC	2400
AGCCCCTGAT	"GCTCTTCGTC	CAGATCATCC	TGATCGACA2	A GACCGGCTTC	CATCCCACTA	2460
CGTGCTCGCT	r cgatgcgatg	TTTCGCTTGG	TGGTCGAATC	GCCACCTACC	CCCATCAACC	2520
GTATGCAGCC	GCCGCATTGC	ATCAGCCATG	ATGGATACTT	TCTCGGCAGG	ACCN ACCMON	2580
GATGACAGGA	GATCCTGCCC	CGGCACTTCG	CCCAATAGCA	GCCAGTCCCT	TCCCCCCCCCC	2640
GTGACAACGT	: CGAGCACAGC	TGCGCAAGGA	ACGCCCGTCC	TGGCCAGCCA	CCATACCCCC	2700
GCLGCCLCGT	CCTGCAGTTC	ATTCAGGGCA	CCGGACAGGT	CGGTCTTGAC	********	
GGGCGCCCCC	' GCGCTGACAG	CCGGAACACG	GCGGCATCAC	AGCAGCCGAT	TCTCTCTCTTCT	2760 2820
GCCCAGTCAT	AGCCGAATAG	CCTCTCCACC	CAAGCGGCCG	GAGAACCTCC	CTCCS MOOR	2820
ICTIGITOAA	TCATGCGAAA	CGATCCTCAT	CCTGTCTCTT	GATCAGATCT	TCAMOCOCOMO	2940
CGCCATCAGA	. TCCTTGGCGG	CAAGAAAGCC	ATCCAGTTTA	CTTTCCACCC	CMMCGGAAAG	3000
LIACCAGAGG	GCGCCCCAGC	TGGCAATTCC	GGTTCGCTTG	CTCTCCATAA	*******	3060
ICIAGCIATO	GCCATGTAAG	CCCACTGCAA	GCTACCTGCT	TTCTCTTTCC	GCTTGCGTTT	3120
TCCCTTGTCC	AGATAGCCCA	GTAGCTGAGA	TTCATCCGGG	GTCAGCACCG	TTTCTCCCC	3120
CTGGCTTTCT	ACGTGTTCCG	CTTCCTTTAG	CAGCCCTTGC	GCCCTGAGTG	CTTGCGGGAC	3240
CGLGWWGCLG	TCAATTCCGC	GTTAAATTTT	TGTTAAATCA	CCAC PARAMAN	TA ACCA ACCA	3300
GCCGAAATCG	GCAAAATCCC	TTATAAATCA	AAAGAATAGC	CCGAGATAGG	COTO & COCOTO	3360
GIICCWGI.I.I.	GGAACAAGAG	TCCACTATTA	AAGAACGTGG	ACTCCAACCT	CARACCCCCC	3420
WWWCCG.LC.L.	ATCAGGGCGA	TGGCGGATCA	GCTTATGCGG	TGTGAAATAC	CCCACACAMO	3480
CGTMAGGAGA	AAATACCGCA	TCAGGCGCTC	TTCCGCTTCC	TCCCTCACTC	XCMCCCMCcc	3540
CICGGICGIT	CGGCTGCGGC	GAGCGGTATC	AGCTCACTCA	AAGGCGGTAA	TACCCOMIANO	3600
CACAGAATCA	GGGGATAACG	CAGGAAAGAA	CATGTGAGCA	AAAGGCCAGC	*****	3660
GWWCCGT.WWW	AAGGCCGCGT	TGCTGGCGTT	TTTCCATAGG	CTCCGCCCCC	CTGACGACGA	3720
I CACAAAAA'I'	CGACGCTCAA	GTCAGAGGTG	GCGAAACCCG	ACACCACTAT	3 3 3 C 3 C 3 C 3 C 3 C 3 C 3 C 3 C 3 C	3780
GGCGTTTCCC	CCTGGAAGCT	CCCTCGTGCG	CTCTCCTGTT	CCGACCCTGC	CGCTTACCGG	3840
ATACCIGICC	GCCTTTCTCC	CTTCGGGAAG	CGTGGCGCTT	TOTOATACION	CNCCCMCMAG	3900
GTATCTCAGT	TCGGTGTAGG	TCGTTCGCTC	CAAGCTGGGC	TGTGTGCACG	*******	3960
I CAGCCCCAC	CGCTGCGCCT	TATCCGGTAA	CTATCGTCTT	GAGTCCAACC	CCCTARCACA	4020
CONCITATO	CCACTGGCAG	CAGCCACTGG	TAACAGGATT	AGCAGAGCCA	CCMAMOMAGO	4080
CGGIGCTACA	GAGTTCTTGA	AGTGGTGGCC	TAACTACGGC	TACACTACAA	CCACACMA	4140
IGGIAICTGC	GUTCTGCTGA	AGCCAGTTAC	CTTCGGAAAA	AGAGTMCCMA	CCMCmmcs mm	4140
COOCHAACAA	ACCACCGCTG	GTAGCGGCGG	TTTTTTTTTTT	GCAAGCACCA	CAMMAGGGG	4200
MONANAAAG	GATCTCAAGA	AGATCCTTTG	ATCTTTTCTA	CTGAACGGTG	ATCCCCACCC	4320
GAATTGCG					G	
						4328

## (2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 5107 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

			K TD 11	O.J.		
GCGGCCGCGG	AATTTCGACT	CTAGGCCATT	GCATACGTTG	TATCTATATC	ATAATATGTA	
CATTTATATT	GGCTCATGTC	CAATATGACC	GCCATGTTGA	CATTOCATONA	MCA COL COM-	60
TTAATAGTAA	TCAATTACGG	CCTCATTACT	TCATAGCCCA	CALIGATIAL	TGACTAGTTA	120
ATAACTTACG	GTB A ATTCCCC	CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	TCATAGCCCA	TATATGGAGT	TCCGCGTTAC	180
AATAATGACC	TIMAMIGGCC	CGCCTGGCTG	ACCGCCCAAC	GACCCCCCCCC	CATTGACGTC	240
CCACMAMMM	TATGTTCCCA	TAGTAACGCC	AATAGGGACT	TTCCATTGAC	GTCAATGGGT	300
GONGIATTIA	CGGTAAACTG	CCCACTTGGC	AGTACATCAA	GTCTN TCNT	mana	360
accect MIT.	GACGTCAATG	ACGGTAAATG	GCCCGCCTCC	Cammamoooo		
or medday.	TITCCTACTT	GGCAGTACAT	CTACCTATION	CTCATCCCT		420
GATGCGGTTT	TGGCAGTACA	CCAATGGGGG	TGGATAGCGG	GICAICGCIA	TTACCATGGT	480
AAGTCTCCAC	CCCATTGACG	TC1 ATTCCC1 C	TOGATAGEGG	TTTGACTCAC	GGGGATTTCC	540
TCCAAAATCT	COMPANDAGO	I CHAIGGGAG	TTTGTTTTGG	CACCAAAATC	AACGGGACTT	600
- constant G1	CGTAATAACC	CCGCCCCGTT	GACGCAAATG	GGCGGTAGGC	CTCTACCCTC	660

GAGGTETTA   ATAGCAGAG   CTCOTTTACT   GARCGETCT   GAGCGCATTC   GAGGGATCC   GAGGGATCC   GAGGGATCC   GAGGGATCC   GAGGGATCC   GAGGGATCC   GAGCGATCC   TATAGCATCA   ATAGCTCATC   GACTATCCAT   ATAGCACCA   GATTATCATC   TATACCACA   GATTATCATC   TATACCACA   GATTATCATC   TATACCACA   GATTATCATC   TATACCACA   GACTACCACC   GACTATCACC   GACCACCACC   GACTATCACC   GACTATCACC   GACTATCACC   GACTATCACC   GACCACACC   GACTATCACC   G	GGAGGTCTAT	ATAACCACAC	CITCCITION OF				
PAGGICACAC   COTTIGGET   TTATICATOR   ANTIACOTOTA   ATTATORCE   ARTORCE   ACTOCOCCE   AC	ACGCTGTTTT	GACCTCCATA	CARCACAGO	GAACCGTCAC	ATCGCCTGG2	GACGCCATCC	720
CCCCGCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	GTGCATTGGA	ACGCCCATTA	CCCCCCCCC	GGACCGATC	AGCCTCCGCC	GCCGGGAACG	780
ATTATTORCE ACTECCETAT TOGGRAGGAT ACTITECTAT ACTARGET 5950 ATTATTORCE ACTIOCCTAT TOGGRAGGAT ACTITECTAT ACTARGET ACTARGET 1140 GACCTOTAT TITTACAGA TOGGRAGGAT ACTITECTAT CHARTCAGA TANCANANA 1140 GACCTOTAT TITTACAGA TOGGRAGGAT ACTITECTAT CHARTCAGA TANCANANA 1140 GACCTOTAT TITTACAGA TOGGRAGGAT CATTORCAGA TOGGRAGGAT GACCTACTAGA TOGGRAGGAT ACTARGAGAT CACCTACTAGA GACTGACAGA ACCTACTAGA GACTGACAGA TOGGRAGGAT ACCTACTAGA GACTGACAGA GACGACAATAT ACCTACTAGA GACTGACAGA TOGGRAGGA TOGGRAGAGA ATTAGACTAC GACCTACTAGA GACTGACAGA TOGGRAGAGA ACCTACTAGA GACTGACAGA TOGGRAGAGA TATAGAGATA TACCTACTAGA GACTGACAGA ACCTACTAGA GACTACACTAT TOGACACTAC CACCTACTAGA GACTACACTAT TOGACACTAC GACCTACTAGA GACTACACTAC GACCTACTAGA GACTACACTAC	TAGGCACACC	CCMMMCCCMC	TOUR TOUR TOUR	GAGTGACGT	AGTACCGCCT	ATAGACTCTA	840
TITEGCACAA CTATCTETAT TIGGTATATG CCAATACTT CTACTTCAGA GACTGACACG   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   1080   108	CCCCCCTCCT	TATCOMAMAC	TTATGCATGC	TATACTGTT	TTGGCTTGGG	GCCTATACAC	900
ACCITITATE THE AGGETATATG   COAPACTOT	ATTATTGACC	ACTICIONAN	GTGATGGTAT	AGCTTAGCCT	T ATAGGTGTGG	GTTATTGACC	960
ACCITICATA THITACAGA TOGGICCECT THATATTA CAATATACA TARACANCAN   1140	TTTTTTTTTT	ACTUCCUTAT	TGGTGACGAT	ACTTTCCATT	ACTAATCCAT	AACATGGCTC	1020
SACTIONAL   THARAGGA   THATHATTHA ALARHAGGA   TARAGACAA   TARAGACAA   TAGCOGTOCT   TAGCOTTCC   CAGACTCCGAG   TAGCATCGG   TAGCATCGG   TAGCATCCGAG   TAGCATCCGAG   TAGCATCCGAG   TAGCATCCGAG   TAGCATCCGAG   TAGCATCCGAG   TAGCATCCGAG   TAGCATCCGAG   TAGCATCCGAGG   TAGCATCCCAC   CACCATTTAC   CACCATTTACA   CACCATTTACAC   TAGCATCCGAGG   TAGCATCCCAC   CACCATTTACAC   CACCATTTACAC   CACCATTTACAC   CACCATTTACAC   CACCATTTACAC   CACCATTTACAC   CACCATTACAC	CACTOTOTOTA	CTATCTCTAT	TGGCTATATG	CCAATACTCT	GTCCTTCAGA	GACTGACACG	
TACOSTITCE GGACATGGGC   TCTTTTTATTA	GUCICIGIAL	TTTTACAGGA	TGGGGTCCAT	ያ ተውጣጥ ይጥቀጥ <b>ፈ</b> ጥጥ	Carammore	M3. M3. M4. M	
CTCCCATCCG	COCCOTC	CGTGCCCGCA	GTTTTTATTA	AACATAGCGT	' CCCRTCTCC	10100	-
CAGACTATOGG   CATGOTCCCT   COCCATGOTC   CAGCACAGO   CACTOGGCOT   1380	TWCQ1Q11CC	GGACATGGGC	TCTTCTCCGG	TAGCGGCGG	COMMOGRA		
SAGETATOTE   TCTGAAAATS   AGCTGAGGA   TAGGGCTCG   CACTGAGGC   CATGGCGGT   AGATGGAGA   CACTGAGGAG   AGATGGAGG   CACTGAGGT   CACTGAGGCA   CAGAGGTATT   CAGAGGGCA   CAGACATAAT   AGCTGAGGGCA   GTGTAGTTT   CAGAGGGCA   CAGACATAAT   AGCTGAGAGA   CAGACATCAT   AGCTGAGGGCA   CAGACATCAT   AGCTGAGAGA   CAGACATCAT   CAGAGGCCAAG   CAGACATCAT   CAGACATCAT   CAGAGGCCAAG   CAGACATCAT   CAGAGGCCAAG   CAGACATCAT   CAGAGGCCAAG   CAGACATCAT   CAGAGGCCAAG   CAGACATCAT   CAGAGAGAGA   CAGACATCAT   CACACACACAC   CACCCACAC   CACCCACACCAC   CACCCACAC   CACCCACAC   CACCCACACCAC   CACCCACACCAC   CACCCACACCAC   CACCCACACCACACCACACACA	GICCCNICCG	TCCAGCGGCT	CATGGTCGCT	- CGGCAGCTCC	THE CONCORD A	G1 000000	
CTTARGACAG   GGGCAGAGA   AGTTGGAGG   AGTGGAGGC   AGTTGGAGGAG   AGTGGAGGAG   AGTGGAGGC   AGTGGAGGAG   AGTGGAGGC   AGTGGAGGC   AGGGTACTC   CGTTGGGGG   GGGGGGAGAT   AGTGGAGGC   AGTGGAGGC   AGGGTACTC   GGGGGGGAGA   AGTGGAGGC   AGGGTACTC   GGGGGGGAGA   AGTGGAGGGC   AGGGTGAGGC   AGGGGGGAGA   AGTGGAGGGC   AGGGTGCGG   AGGGGGAGA   AGGGGTGCG   AGGGTGCGG   AGGGCGAGA   AGGGGTGCG   AGGGGGGAGA   AGGGCCACCC   AGGATGCAGG   AGGGCGAGA   AGGGCCACCC   AGGATGCACC   AGGATGCAGG   AGAGTGCAGG   AGGGCGAGA   AGGGCCACCC   AGGATGCACC   AGGAGGGCCAG   AGGGCCACCC   AGGAGGGCCAG   AGGGCCAGAC   AGGGCCACCC   AGGAGGGCCAG   AGGGCCAGAC   AGGGCCAGCC   AGGAGGGCCAG   AGGGCCAGAC   AGGGCCAGCC   AGGAGGGCCAG   AGGGCCAGCC   AGGAGGGCCAG   AGGGCCAGAC   AGGGCCAGCC   AGGAGGGCCAG   AGGAGGGCCC   AGCACCACCC   ACCACCACCC   ACCACCACC   ACCACCACCC   ACCACCACCC   ACCACCACC   ACCACCACCC   ACC	CHORCITAGG	CACAGCACAA	TGCCCACCAC	CACCAGTGTG	CCCCACAAAA	GG0000	
CASTILLATION   CONTINUED   C	PIDIATOOON	TUTGAAAATG	AGCTCGGAGA	一	ACCMCC ACCC	10100-	
STIGNTECT   GCTOTTGCGST   GCTOTTTACC   GTGCATGCCA   GTTCCTTTCC   GTCCTTTCC   GTGCTTTCC   GTGCTTCTT   GTGCATGCCA   GTTCCTTCCC   GTTCCTTTCC   GTGCTCTCC   GTGCTTCTC   GTGCTCTCCC   GCTCATCATCA   GTCCTCCCC   GCTCATCATCA   GTCCTCCCC   GCTCATCATCA   GTCCTCCCCC   GTCCTCATCA   GTCCTCCCCC   GTCCATCATCA   GTCCTCCCCC   GTCCATCATCA   GTCCTCCCCC   GTCCATCATCA   GTCCTCCCCC   GTCCATCATCA   GTCCTCCCCC   GTCCATCATCA   GTCCTCCCCC   GTGCATCATCA   GTCCATCACCC   GCCCCACCACCC   GTCCATCACCC   GTCCATCCACC   GTCCATCCACCC   GTCCATCCACC   GTCCATCCACC   GTCCATCACCC   GTCCATCCACC   GTCCATCCACC   GTCCATCACCC   G	CTIMOGCAG	CGGCAGAAGA	AGATGCAGGC	AGCTGAGTTG	നന്നാന മന്ദ്രാനം	3 603 3 604 604	
ATGGGTCTTT   TOTGGGGCC   CAGACMTAAT   AGCTGACAGA   CTAAGACTT   CAGACTCGAG   CAAGTCTAGA   TOTGGCCTCT   TOTGGCGCCAC   GGCGCGGCA   GGGGGGGCAC   CGAAGTCCTC   GCCTCACTCT   TOTGCCCCCT   TOTGCTCCCTC   TOTGCTCCCCT   TOTGCTCCCCT   TOTGCCCCCC   GAAGGCCCAGA   GGCGCCAGAC   GCCCCACCAC   1800   GCCCCACCAC   CACACTCCCCC   GAAGCCCAGAC   GAAGCCCAGA   GGTCCCCCCC   GAAGCCCAGAC   GCCCCACCAC   CACACCCCCCCCCC	GUGGITACIC	CCGTTGCGGT	GCTGTTAACG	GTGGAGGGCA	CTCTTCTTCTC	3.553.55	
TOGOTGAGGG   ACCCGGGCA   GGGGGGGAGA   TGGGGGTGCA   CAAATGTCAG   CAAAGTCAG   GGCTGAGCTT   TGGTCTGGT   TGGTCTGGT   TGGGGCTCC   TGGGGCTCC   AGTCCCGGGG   GCCCGCGCAG   GGCCGGCGAG   GGCCTGGCTT   TGGAGCCCAG   GGCTCACCCAC   T800   GCCTCACCCAC   TGTCATGGG   GGCTGGTT   TGGAGGCGAG   GGCAGAGGCG   GGCAGAGGCC   TGTCATGGG   GGCAGAGGCT   TGGAGGCGAG   GGCAGAGGCC   TGTCATGCC   GGCAGAGGCC   TGGAGGCGC   TGGAGGCGC   TGGAGGCCG   TGGAGGCCG   TGGAGGCCG   TGGCAGCAGC   CTCCCAGACC   TCTCATGCG   GGCCTGCGCC   TGGAGGCGC   TGGCAGCACC   CTCCCGGGCC   TGCCAGCCC   CCCAGGAGGC   CCCAGGAGGC   CCCAGGAGGC   CCCAGGAGCC   CCCCAGGACC   CCCCCACACC   CCCCCACCC   CCCCCACCC   CCCCCACCC   CCCCCACCC   CCCCCACCC   CCCCCCACCC   CCCCCACCC   C	GIIGCIGCCG	CGCGCGCCAC	CAGACATAAT	AGCTGACAGA	CTAACACACA		
GOCTIONISMS   GOCCEGO	MIGGGICITT	TUTGCAGTCA	CCGTCGTCGA	ייייי ב בטב ביייי	CACACMOORG	@X > ===	
SCCTIONTECT  TOACAGCCCA   TOCTOTOCO   TOACTCCACCA   TOACTCCACAC   TOACTCCACACACACACACACACACACACACACACACACA	TCGCTGAGGG	ACCCCGGCCA	GGCGCGGAGA	TGGGGGTGCA	CAGACTCGAG	CAAGTCTAGG	1680
ARGITCACAST   GAGGCTATTC   GAAAGCTGCA   GATTATCACC   GTCCCAGCA   1920	GGCTTCTCCT	GTCTCTCGTG	TOGOTOCOTO	TEGECCTCCC	CGMATGTCCT	GCCTGGCTGT	1740
1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920   1920	GCCTCATCTG	TGACAGCCGA	GTCCTCGAGA	CCERCCOCC	AGTCCCGGGC	GCCCCACCAC	1800
GGCAGGGCT GGCCCTGCTC TCGAGAGGCT GGCAGGCCT GTGCAGCCC CTGCCAGCC TTTCCCAGCC TTTCCAGCCC CTGCAGCCC TTTCCAGCCC CTGCAGCCC CTGCAGCCC CTGCAGCCC CTGCAGCCC CTGCAGCCC CTGCAGCCC CTGCCCCCC ACCCTCCCCC ACCCTCCCCC ACCCTCCCCCC ACCCTCCCCC ACCCTCCCCC ACCCTCCCCC ACCCTCCCCC CTGCCCCCCC CTGCCCCCCC CTGCCCCCC CTGCCCCC CTGCCCCCC CTGCCCCC CTGCCCCCC CTGCCCCC CTGC	ATGTCACGAT	GGGCTGTTCC	GAAAGCTCCA	COMMONATOR	GGAGGCCAAG	GAGGCCGAGA	1860
CTTCCCAGC  TTTGGAGCC  TTTGGAGCC  TTTGGAGCC  TTTGGAGCC  GCATCACCA  CAGAGGAGGC  GCATCACCAC  GCATCACCAC  GCATCACCAC  CTCCGGCTGC  TCCCCGGCTGC  TCCCCGGGGGA  ACCATCACTG  ACCCAAATTC  CTCCCGGGGGA  ACCTCACCTC	CCAAAGTTAA	CTTCTATCCC	TOGARCACCA	GCTTGAATGA	GAATATCACC	GTCCCAGACA	1920
GATTACACCAC TYTUGAGUCCE CTGCAGCTGC ACATGGATAA AGCCATCAGT GGCTTTCGCA GCATCACCAC TCCCCTTCGGA ACCTTCACTG CTGACACTTT CTCCAAACTC CTCAGATTCT ACTCCAATTT CTCCCGGGGA AAGCTGAAGC TGTACACGG GGAGGCCTGC AGGAGAGGG ACAGATGAGC AGGTGGGTC ACCTGGACAC TCCCTCACCA ACCTGCCTG TGCCACACCC TCCCTCACCA CTCCCGAACC CTCCCTCAGCA AACCTGCCTG TGCCACACCC TCCCTCACCA CTCCCGAACC CCATCAGGG GCTCTCAGCT AACGGCCAGC 2460 AGCACACACT TGGAGTCCAC TCCCGAACC CCATCAGGG GCTCTCAGCT AACGGCCAGC 2460 AGCACACACT TGGAGTCCAC TCCCGAACC CCATCAGGG GCTCTCAGCT AACGGCCAGC 2460 AGCACACACT TGGAGTCCAC TCCCGAACC CCCCGTGCCT TCCTTCAGCT AACGGCCAGC 2460 AGCACTCACT TGGAGTCCAC TAGCGCTTGA GCCCCGTGCCT TCCTTGACCC TGGAGGTGC 2580 TAGCACGGGGG GTGGGGGCA GACATCCAC TGCGAGGGGG GCCCTGACCA AACGCCCAC CTTGCCTTTC TTTGCCCCT CCCCGTGCCT TCCTTGACCC TGGAGGGGG 2580 TAGCACGGGGGG GGGGGGCA GACATCACA GGGACACAAG GGGGAGGACAA ACCCCCAC TGGAGGGGG GGACAGCAAG GGGACACAAG GGGGAGACAAA ACCCCTAAACAC CCGAAGCCCA ACCTTTCATA GAAGGCGGG ATCATCCAGCA TGGAGACCCA ACCTTCAACAC ACCCTGAAGCACA GGCGCTGCCT TCCTCAACAC ACCCTTAAACAC ACCCTAAACAC ACCAGAAAG ACCCTAAACAC GGCGTCACCT GTCAACACA GGCGACACAC GGACACACA GGCGCCACAC GAATACAA GCGCACACAC ACCTTTCATA GAAGGCGGC TTCCACCAA ACCCTAAACAC ACCCTAAACAC ACCCTAAACAC ACCCCAACAC ACCCTAACAC ACCCTAACAC ACCCTAACAC ACCCCAACAC ACCCTACACAC ACCCTAACAC ACCCCAACAC ACCCTACACAC ACCCCAACACC ACCCTACACAC ACCCCAACACC ACCCCAACAC ACCCCAACAC ACCCCAACAC ACCCCCAACAC ACCCCCAACAC ACCCCCAACAC ACCCCCAACAC ACCCCCAACAC AC	GGCAGGGCCT	GGCCCTCCTC	TGGAAGAGGA	TGGAGGTCGG	GCAGCAGGCT	GTAGAAGTCT	1980
CCTCGGCTGC TCCACTCGG GCGCTGAGAG CCAGAGAGC CATCTCCCTC CCACATCCTG ACTCCAAACTC TCCCAAACTC TCCCAAACTC TCCCAAACTC TCCCAAACTC TCCCAAACTC TCCCAAACTC TCCCAAACTC TCCCAAACTC ACTGCCAAACACACACACACACACACACACACACACACAC	CTTCCCAGCC	TTTCCACCCC	TCAGAAGCTG	TCCTGCGGGG	CCAGGCCGTG	TTGGCCAACT	2040
CATCCATTT         CTCACTCGGA         AGCTGAACT         CTTGCAACTT         CTTGCAACTC         2220           ACTCCAACTT         CTCGGGGA         AGCTGGGCA         CTCACCAGCG         CAGGAGGGG         2280           ACACACACC         TCCCTCACCA         CTCCCGAACC         CTCCCGAACC         CTCCCGAACC         CTCCCGACCA         AACTCCACG         2400           AGCACAACTC         TGAGATCCAC         TACGCGTTAG         AGCTCGCTGA         AACTCTCAGC         2460           AGCACAACTC         TGAGATCCAC         TACGCGTTAG         AGCTCGCTGA         TCCATCCCCAC         CATCTCCACCAC         TCCCTTTCCT         AACTCTCAGCA         ACCCTTGACCT         TCCATCCCCAC         TCCATCTCACCAC         TCCCGTTGACCT         TCATTGACCC         AGCAGACACACACACACACACACACACACACACACACAC	GCATCACCAC	TITCGAGGGG	CTGCAGCTGC	ACATGGATAA	AGCCATCAGT	GGCCTTCGCA	2100
ACTECASTT         CUCCTEGGGGA         AGGTGAAGT         CTGCAAGTT         CTGCAGGGG         2220           ACACACACC         CACCTEGGGAA         AGCTGAGGC         2280           ACAGATGACC         AGCTGAGCA         ATCCACCACC         CTCCTCACCA         ACCTCGCTCACCA         ACCTCCCTCACCA         ACCTCCACTCA         CACTCCACTCA         CACTCCACCAC         CCCCGAACC         CTCCCGACCAT         ACCACTCACTCA         AACCTCCACTCACCA         CACTCCCACCAC         CACTCCCACCAC         CCCCGTGCCT         TCCTTGCCTC         2460           AGCACACCT         TGGAATCCAC         TACGCGTTAG         AGCTCGCTCACCAC         GACTCCCCCAC         CACTCCCACACT         TCCTTTCCT         TATAGTTGCCC         CCCCTTTCCT         ATAAAAATGA         GACACCACAC         TCCGATTGTC         TCAGTTGAGCAC         TCCGATTGTC         TCAGTTGAGCAC         ACCTTTATT         CTGGAAGCCA         ACCTTCACACA         AGCGCACCACA         TCCGATTGTC         CTGAAGCCCA         ACCTTCACACA         ACCTCACACACC         CAGGACACCC         ACTCACACCACA         ACGCCATACACCA         ACCTCACACACACCACACACACACACACACACACACACA	CCTCCCCTCC	TCTGCTTCGG	GCGCTGGGAG	CCCAGGAAGC	CATCTCCCTC	CCAGATGCGG	2160
ACCIDATION         COTTOGGGGA         AGCTGGACC         TECTACAGGG         GGAGGGGCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	ACTCOOLIGC	TCCACTCCGA	ACCATCACTG	CTGACACTTT	CTGCAAACTC	TTCCGAGTCT	
TOCACACCC   TCCCTCACA   TCCCCGAAC   TCCCCCACA   ACACTGCCTG   2340	MCICCMMIII	CCTCCGGGGA	AAGCTGAAGC	TGTACACGGG	CORCOOMOO	100101	
CRETECCATE GRACETECAS GRACACTECAS GRACACTECAS GRACACTECAS GRACACTECAS AGCACAACTE AGCACACACTE AGCACACACTE AGCACACACTE AGCACACACT TAGTTCCAS CCATCEGTTG TTTGCCCCT CCCCGTGCTT TCTTGCCCT CCCCGTGCTT TCTTGCCCCT CCCCGTGCTT TCTTGCCCT TCGATGCTC TCGATCGTTC TTTGGCGGGCA GGGCGCAGAG TCGATCCACT TCGGGGGGGAG AACTCCAGCA TGGAGCGAG TCGATCCACT TAGGGGGGGAG AACTCCAGCA TGGAGCGAG TCGCGTCCC GGAAACGATT CCCAAGCCCA ACCTTCATC TGGAGGGAG AACTCCAGCA TGGAGCGAG TGGCGAAGA ACCTCCAGCA TGGAGCGCA TGGCGAGCG CGGCGCCCC GGAAACGATT CCCAAGCCCA ACCTTCATA AACTCCAGCA TGGAGGGAG ACCTTCATA TAGCAGGAGA CCGGCGCCCC CGGCGTCCC GAAACGATT CCCAAGCCCA ACCTTCATA ACCTTAAAC ACCGTAAACA ACCGTAAACA ACCGTAAACA ACCGTAAACA ACCGTAAACA CGGCACTTT CCCAAGCACC CGGCGCCCACA CGCACACCAC TCCAGCACCA TCCACACACCA TCCACACACCA ACCCTCACAC CGCACCACCA CGCCACCACC CGCACCACC CGCCACCACC CGCCACCACC CGCCACCACC CGCCACCACC CGCACCACC CGCCACCACC CGCCCCCC CCGCCACCACC CCCCCCCC	ACHONIGACC .	AGGTGCGTCC	AGCTGGGCAC	ATCCACCACC	がたたたかのみのです	1010man	-
AGCACACTC TGAGATCAC TGCCAGCAT TACGCCTCAC GGCCCCCAC CTGGCCCTCAC CTGTCCCTTC CTGAGGTCAC CACTCCACT GTCCTTTCCT ATTGCCCCTC CCCCGTGCCT TCCTTGACCC TGGAAGGTGC CACTCCACT GTCCTTTCCT AATAAAATGA GAAATTGCA TCGCATTGCT TGGAAGGTGC CACTCCACT GTCCTTTCCT AATAAAATGA GAAATTGCA TCGCATTGCT TGGAAGGTGC CACTCCACTATTCATT CTGGGGGGTG GGGTGGGGCA GGACACCAAG GGGGAGGATT GGGAAGCAAA CACTCCACCAC TGGAAGCACA ACCTTCATA GAAGGCGGC GTGGAATCCACC CACAGCCCC GCGCTGGAGG ATCATCCACC CACAGCCCC GAAAACGAAT CCGCAAGCCCA ACCTTCATA GAAGGCGGC GTGGAATCGA 2820 ACCTTCCACAAAAACACAC TCGTCAAAAA GGGGAAGACAA GGCGATACGA ACCTTCACAC AGAGTCCCCC CACAGCCCC ACCTTCACACAAAAACACAC CCCACAAAAAACACAC CACACAAAAACACAC CACACAAAAAA	* OCCUCACOC	TCCCTCACCA	CTCCCGAACC	CCATCGAGGG	COTOTOROROR	* * * * * * * * * * * * * * * * * * * *	
ACCARACTO         TRAGECCATC         TAGGCGTTAGA         AGCTCGCTGG         TAGGCGTCC         2520           CACTCCCACT         GTCTTTCTT         TATTGCCCCTC         CCCCGTGCCT         TCCTTGACCT         TGGAAGGTGC         2580           CACTCCCACT         GTCTTTCTT         AATAAAATGA         GGAAATGCAT         TGGAATGCT         TGGAAGAGAA         2700           TAGCAGGGGG         GTGGGCGAA         AACTCCAGCA         TGGAATCCC         GGGAGGAGA         ATCATCCAGC         2760           CAGCGTCCCG         GTGGAAGCCA         CCGAAGCCCA         CTCTACAACCC         GGCGTGGCT         GGCGAAGCCA         ATCATCCAGCA         AGAGGCGGC         AGAGGCGCCA         ATCATCCAGCA         GAGGCGCCA         ACTCAAAACC         AGAGGAGAGC         GGCGATCACC         AGCGGAAACCA         AGAGGAGAGC         GGCACCCA         ACTCACAACCA         AGACGGAACCA         AGCTCATCCA         CAATACCACCA         AGACGCACCA         AGACCGCCACACCA         AGCCGGCCACCACCA         AGCCGGCCACCACACACACACACACACACACACACACAC	CIGICCCAIG	GACACTCCAG	TGCCAGCAAT	GACATCTCAG	GGGGGAGAGA	3 3 CEC	
CACTCCCACT GTCCTTTCCT AATAAAATGA GGAATTGCA TCGCATTGCT TCGTGAGGTG 2640 TCATTCTATT CTGGGGGGG GGGGGGCA GGACAGCAAG GGGAGGACAA GGGAGGCGCAGGGGGGGG	AGCACAMCTC .	TGAGATCCAC	TACGCGTTAG	AGCTCGCTGA	TO LOCOTRODA	amana a a a a a a a a a a a a a a a a a	
TCATTCTATT CTGGGGGGTG GGGTGGGCA GGACAGCAAG GGGGAGGATG ATCATCCAGC TAGCAGGGG GTGGGCAAG AACTCCAGCA TAGAATCCCG GGGGAGGATG ATCATCCAGC AATCCAGCA AACTCCAGCA ACCTTCATA GAAGGCGC GGAAGACAAG GGGGAGGATG ATCATCCAGC 2760 AAAACGACAT TGGCAAGACA GGCGATAGAA GCGCACACC AGAGCACCC AGAGTCCACC AGAGCACCC AGACCCCACCC AGACCCCACCC AGACCACCC AGACCACCC AGACCACCC AGCCACCC ACCACCCAC	THG T TOCCAG	CCATCTGTTG	TTTGCCCCTC	CCCCGTGCCT	<b>ምሮሮሞምር አ</b> ሮሮሮ	mcca a comes	
TAGCAGGGGG GTGGGCGAAG ARCTCCAGCA TAGAGTCCCC CGCCTGGAGG ATCATCCAGC 2760 CGGCGTCCCG GAAAACGATT CCGAAGCCCA ACCTTCATA GAAGCCCG GTGGAATCGA 2820 ARCTCGTGA TGGCAGGATG GGCGTCGTT GGTCGGTCAT TTCGAACCC AGAGTCCCC 2880 TCAGAAGAAC TCGTCAAGAA GGCGATAGAA GGCGATCGC TTCGCGGCAA AGAGTCCCCG 2880 ACCGTAAAAC ACGAGGAAGC GGTCAACCCA TTCGCGGCCA AGCTCTTCAG GAAGCCGAACCC CATAGTCCAG GGTAGCCAA ACGCGCAACCC AGCCGCAACCC CAGCCCAACCC CAGCGCAAC CATCGAGAAAAG CGGCCATTT CCACCATGAT ATTCGCGGCCA AGCCGGCAAC AGCCGGCAAC AGCCGGCAAC AGCCGGCAAC CATCGGTCAC AGCCGCACACCC CAGCGCAACCC CAGCGCAACCA CAGCGCAACCC CAGCGCAACCA CAGCGCAACCC CAGCGCAACCA CAGCGCAACCA CAGCGCAACCA CAGCGCAACCA CAGCGCAACAC CAGCGAACAC CAGCGAACAC CAGCGAACAC CAGCGAACAC CAGCGCAACCA CAGCGCACACC CAACCAGCACC CAACCAGACAC CAGCGCACACC CAACCAGCACC CAACCAGCACC CAACCAGCACC CAACCAGCACC CAACCAGCACC CAACCAGCACC CAACCAGCACC CAACCAGCACC CAACCAGCACC CACCACACCA CACCAACCA	CHCICCCMCI	GICCLTLICCT	AATAAAATGA	GGAAATTGCA	TCCC A TYPOTO	MC 3 CM 3 COmo	
CGGGTCCCG GAAACGATT CCGAAGCCCA ACCTTTCATA AAAGGCGGG GTGAATCGA 2820 AATCTCGTGA TGGCAGGTT GGCGTCGCTT GGTCGGTCAT TTCGAACCCC AGAGTCCCG 2880 ACCGTAAAGC ACGAGAAGC GGCGATAGAA GGCGATAGCA GGCGAATCGC TGCGAATCGG GAAGGCGGGAT CAAAAGAAAGCCCA ACCTTCAAGAA ACCTCTCAG CAAAAGAACACTC TCCAGAAAAG CGCCAATGAA GGCGAATGAA ACCTCTCAG ACCCC AGCGCAACCC AGCGCAACCC AGCGCAACCC AGCGCACCC AGCGCACACCC AGCGCACCC AGCGCACACC AGCGCACCC AGCGCACCC AGCGCACCC AGCGCACCC AGCGCACCC AGCGCACCC AGCCACCCAC	TCMITCIMIT (	CIGGGGGTG	GGGTGGGGCA	GGACAGCAAG	GGGGAGGAmm	CCCANOXONA	
CARTICLEGIA         CARAMAGART         COCARAGCCCA         ACCITTCATA         GARGGGGGG         GTGGAATCGA         2820           AATCTCGTBA         TGGCAGGTTG         GGCGTCGCTT         GGTCGGTCAT         TTCGAACCCC         ACAGTCCCGC         2880           ACCGTAAAGC         ACGAGGAAGC         GGCGATAGAA         GGCGATACACA         AGCTCTTCAG         CAATATCACG         3000           GTTAGCCAA         GCGCACACCC         AGCCGCCCA         AGCCGCCCA         AGTCGATGAA         3060           GACGAGATCC         CCGCACACCAC         AGCCGCCCC         AGTCGATGAA         3060         3000           GACGAGATCC         CCGCCACATTC         CCACACATGAT         ATTCGGCCAC         AGCCGGCCCC         CATGGGTCAC         3120           GAGCCCCTGA         TGCTCTTCGT         CCACACTACAT         CTTGAGCCTG         CCTGACACGC         CCTGACACGC         CCTGACCGCGC         CCTCACCGAAC         CAACCGCCGT         CCTCACCGACC         CAACCGCCGCT         CCTCCGCATG         GCCCCAATAGC         CAGCACAGCC         CCGGCATTCC         CCGGCATTCC         CCGGCATTCC         CCGGCATTCC         CCGGCACTCC         TTCCCGCTTC         3420           CGCGCACTCC         TCCGCCTGCAC         CTGCCCATACAC         ACCGGCCCTC         TTCCCGCTTC         TTCCCGCTTC         CCAACAGCCC         CTTGCCAACC	TUGCUGGGG	GTGGGCGAAG	AACTCCAGCA	TGAGATCCCC	GCGCTGCACC	A MCAMOON OO	
TCAGAAGAA TCGTCAAGAA GGCGATAGAA GGCGATAGAA GCGCATAGAC ACGGTAAAGA ACGGCAAAAGA ACGGCAAAAGA ACGGCAAAAGA ACGGCAAAAGA CAGAGAAAGAC GGTCAGCCCA TTCGCCGCCA AGCTCTTCAG CAATACACAG 3000 TCCAGAAAAAA CGGCCATTT CCACCATGAT ATTCGCCGCCA AGCCATCAC CAATACACAG 3000 ACGGAAAAAA CGGCCATTT CCAGACACAC AGCCGCAACACC CATGGTCAC CAATACACAG 3120 ACGCCCTGA TGCTCTTCGT CCAGAACACT CCAGATCAT CTGAGCCCT TCGAGCCAC CATGGGCAC CATGGGCACAC CAGACAGGTT CATCCAGCCAC AGACCAGCTAC TTCCCGCAGT 3240 ACGCCAGCCAC AGACCAGCCC TTCCCGCATC CATCCAGCACAC CAGACAACAC CAGACAAGAC CACCAATAGC ACCAATAGCC ACCAATAGCC ACCAATAGCC ACCAATAGCAC ACCAATAGCC CAAAAAAAAAA	COCCE LCCC	GAAAACGATT	CCGAAGCCCA	ACCTTTCATA	GAAGGGGGGG	CMCC3 3 mmcs	
ACCGTAAAGA GGCGATAGAA GGCGATGCCC TTCGCCGCCA AGCTGATCAC CAATATCACG GATAGCAAC GGTTAGCCAC GGTTAGCCAC GGTTAGCCAC GGTTAGCCAC GATAGCGGTC CGCCACACCC AGCCGGCACCC CAATGCGATGAA 3060 AGTCGAAAAA GCGGCGCCACCC AGCCGCCACCC AGCCGGCACCC CAATGCGATGAA 3060 AGTCGAAAAAA GCGCGCCACCC AGCCGCCACCC AGCCGCCACCC CAATGCGATGAA 3060 AGTCGAAAAAACCGCCACCC AGCCGCACCC AGCCGCACCC CAATGGCACCC CAATGGATAAAAACCGCC TTCGCCGTCG GCAATGCACCC AGCCGCACCC AGCCGCACCC CAATGGACACC CAATGGACACC CAATGGATCAC CCACAACACCC CAATGGACCCC CAATGGACCCC CAATGGACCCC CCAATGACC CCAGATCACC CCAGATCACCC CCAGATCACC CCAGATCACC CCAGATCACC AGACCGGCTT CCATCCGACT 3240 ACGTGACAACG AGATCCTGCC CCGGCACTTC GCCCAATAGC AGCCAGTCAC CCAGATGAC AGACCAGCTC TTCCCGCTTC 3420 ACGTGACAACG TCCGCCACCC TCCGCCACCC ACGATCACCC ACGATACCC ACCGACCACC ACGATAGCC ACGATAGCC ACGATAGCC ACCGACCACC TCCGCCACCC TCCGCCTTC ACCCGACCC TCCGCCACCC ACCGACCCC ACGATAGCC ACCGACCCC TCCGCCTTC ACCCGACCC TCCGCCACCC ACCGACCCC ACCCACCC	MATCICGIGA .	1GGCAGGTTG	GGCGTCGCTT	GGTCGGTCAT	THECONNECCO	3 C 3 C 0 C 0 C 0 C 0 C 0 C 0 C 0 C 0 C	
GGTAGCCAAC GCTATGTCT GATAGCGGTC CGCCACACCC AGCCGCCACCC AGCCGGCCACCC AGCCGCCACCC AGCCGGCCACCC AGCCGCCACCC AGCCGCCACCC AGCCGCCACCC AGCCGCCACCC AGCCGCCACCC AGCCGCCACCC AGCCGCCACCC CGCCACCCC CTGGCGCC CTTGAGCCACC CAGCGCCACCC CAGCGCCACCC CTGAGCACCC CTGAGCACCC CTGAGCCACCC CTGACCCACCC CTGACCCACCC CTGACCCACCC CTGATCGACA AGCCCGGCTT CCATCCGAGT 3240 ACCGTCCCCCC CCGCCACTC CCACCCACCC AGCCACCCC CCGCCCACCC CCGCCCACCC CCGCCCACCC CCGCCCACCC AGCCACCC CTGATGCACA AGCCCGGCTT CCCTCCACC AGCCACCC CCGCCCACTC CCCGCCACTC CCGCCACCC AGCCAGCTC CCCGCCACCC CCGCCCACCC CCGCCCACCC CCGCCCACCC CCGCCCACCC CCCGCCACCC CCGCCCACCC CCCGCCACCC CCCCCCCC	I CAGAAGAAC .	PCGTCAAGAA	GGCGATAGAA	GGCGATGCGC	TGCGAATICCC	CACCCCCCA	
TCCAGAAAAG CGGCCATTT CCACCATGAT ATTCGGCAG CAGGCATCC CAGCCATGAT CCACCATGAT ATTCGGCAGC CAGGCATCC CAGCCATGAT CCACCATGAT CCACCATGC CCCCACTCC CCGCCACTC CCCCCACTCC CCCCCACTCC CCCCCACTCC CCCCCACTCC CCCCCACTCC CCCCCCACTC CCCCCACTCC CCCCCCCC	ACCGIAAAGC ,	AUGAGGAAGC	GGTCAGCCCA	TTCGCCGCCA	ACCOCOOCAC	03 3 m 3 m 0 3 m 0	
GACGAGATCC TCGCCGTCGG GCATGCGCCC CTTGAGCCTG GCGAACACTT CGGCTGGCGC 3180 GAGCCCCTGA TGCTCTCGT CCAGATCATC CTGATCCACA AGACCGGCTT CCATCCGAGT 3240 ACGTGCTCGC TCGATGCGAT GTTTCGCTTG GTGGTCGAAT GGGCAGGTAG CCGGATCAAG 3300 CGTATGCAGC CGCCGATTG CATCAGCCAT GATGGAACA GGCCAGGTAG CCGGATCAAG 3300 AGATGACAGG AGATCCTGCC CCGGCACTTC GCCCAATAGC AGCCAGTCCC TTCCCGCTTC 3420 AGTGACAACG TCGAGCACAG CTGCGCAAGG ACCGGACAGGTCC TTCCCGCTTC GCGCGCCCCC TCGCGCAGAG ACCGGCACCC ACGGACCAG TCGGCCCCCC TCGCGCAAG GCCGGACACG TCGGCCCCC TCGCCGAACA GCCGGCACCC ACGGACACG TCGCCCAACACG ACCGGCCCC TCGCCGAACAC GCCGGACACG TCGGCCAACAC GCCGGCACCC ACGAAAAAGAAC 3540 CGGCCCCCC TCCGCTGACA GCCCGAACAC GCCGGACACG TCGGTCTTGA CAAAAAGAAC 3540 ATCTTGTTCA ATCATGCGAA ACGATCCTCA TCCTGTCTCT TGATCAGATC TTGTCTGTTG 3600 ATCTTGCAACAA ATCCTTGGCG GCAAGAAAGC CATCCAGTTT TGATCAGATC TTGATCCCCT 3720 CTTACCAGAG GCCCCCAG CTGGCAATTC CGGTTCCTCT TGATCAGATC TTGATCCCCT 3720 CTTACCAGAG GCCCCCAG CTGGCAATTC CGGTTCCCTT TGATCAGATC TTGATCCCCT 3780 CTTACCAGAG GCCCCCAG CTGGCAATTC CGGTTCCCTT TGATCAGATC TTGATCCCCT 3780 CTTACCAGAG GCCCCCAG CTGGCAATTC CGGTTCCCTT TGATCACACG GCTTCCCAAC 3840 ACTCGCTTTC CAGATAGCCC AGTAGCTGC ACCTCCTT TTCTCTTTTC CGCTTCCAAC 3840 ACTCGCTTTC TACGTGTTCC GCTTCCTTTA GCACCACCC GTTTCCCATC ACTTTCTTTTC CGCTTTCCTTTC	GGINGCCAMC (	GUTATGTCCT	GATAGCGGTC	CGCCACACCC	ACCCCCCCA	1.000000000	
GAGCCCTGA TGCTCTTCGT CCAGATCATC CTGATGCACA AGACCGGCTT CCGTATGCAC CCGGCCCTGA TGCTCTTCGT CCGAGTCACC CTGATCGACA AGACCGGCTT CCGCCGCATTG CATCAGCCAT AGATGACAGG AGATCCTGCC CCGGCACTTC CCGGCACATG CCTACCGCAC AGATGACAACG AGATCCTGCC CCGGCACTTC CCGGCACATG CCCCCAATAGC AACGCCCGTC CCGGCACATG CCCCAATAGC AACGCCCGTC CCGGCACATG CCCCAATAGC CCGCCCAATAGC CCGGCCCCC TCCGCCACAG CCCCAATAGC CCGGCCCCC TCCCGCACAC CCGGCACACG CCCCAATAGC CCGGCCCCC TCCCGCACAC CCCGGCACAC CCCGGCCCCC TCCCCCACC CCGGCCCCC TCCCCCACC CCCGGCACAC CCCAGCCCCC CCCCCCCC CCCCCCCC	+ ccuguuuu	CGGCCATTTT	CCACCATGAT	ATTCGGCAAG	CAGGCATCCC	CAMCOCMOAG	
ACGTCCTCGA TGCTCTCGT CCAGATCATC CCAGATCATC CCGTATCCAC CCGCGCATTG CCATCCCCCC CGCCGCATTG CATCACCCAC AGATGCACAC AGATGCACC CGCCGCATTG CATCACCCAC AGATGCACAC AGATGCACC CCGCGCATTG CATCACCCAC AGATGCACC CCGCCCCATC CCGCCCCATC CCGCCCCATC CCGCCCCATC CCGCCCCCCC CCGCCCCCC CCGCCCCCC CCCCCCC	OUCOMONICC .	redecerege	GCATGCGCGC	CTTGAGCCTG	GCGAACACTO	CCCCMCCCCC	
CGTATGCAGC CGCCGCATTG CATTCAGCCAT GATGGATACT CATGACCAT AGATGACAACG AGATCCTGCC CCGCGCATTC CCGCGCATTC CCGCGCATTC CCGCGCATTC CCGCGCATTC CCGCGCATTC CCGCGCATTC CCGCGCATTC CCGCGCATTC CCGCGCATCC CCGCGCATCC CCGCGCATCC CCGCGCATCC CCGCGCATCC CCGCGCACTC CCGCCCCCC CCGCGCCCC CCGCGCACCC CCGCGCCCC CCGCGCCCC CCGCGCCCC CCGCCCCCC	GAGCCCCTGA 1	TGCTCTTCGT	CCAGATCATC	CTGATCGACA	ACACCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	CGGCTGGCGC	
AGATGACAGG AGATCCTGCC CCGGCACTTC GCCCAATAGC AGCCAGTCC TTCCCGCTTC 3420 AGATGACAAGG TCGAGCACAG CTGCGCAAGG AACGCCCGTC GTGGCCAGCC ACGATAGCCG 3480 CGCTGCCTCG TCCTGCAGTT CATTCAGGGC GCCGACACAG TCGGCCAGCC ACGATAGCCG ACGGCCCCC TGCGCCAACA ACGGCCAGCC ACGATAGCCG TTCCCGCTTC GCCGCACACA ACGGCCAGCC ACGATAGCCG ACGGCCCCC TGCCCACACACACACACACACACACACACACACACACACA	ACGTGCTCGC	TCGATGCGAT	GTTTCCCTTC	GTGGTCGA AM	ACACCGGCTT.	CCATCCGAGT	
AGTGACAAGG AGATCCTGCC CCGGCACTTC GCCCAATAGC AGCCAGTCC TTCCCGCTTC 3420 AGTGACAACG TCGAGCACAG CTGCGCAAGG AACGCCCGTC GTGGCCAGCC ACGATAGCCG 3480 CGGCGCCCC TCCTGCAGTT CATTCAGGGC ACCGGACAGG TCGGTCTTGA CAAAAAGAAC 3540 TGCCCAGTCA TAGCCGAATA GCCTCTCCAC CCAAGCGGCC GGAGCACCG TTGTCTGTTG 3600 TTCCTTGTTCA ATCATGCGAA ACGATCCCAC CCAAGCGGCC GGAGAACCTG CGTGCAATCC GCACCACAGA ACCGCCCACAGA TCCTTGCCAC CCAAGCGGCC GGAGAACCTG CGTGCAATCC GCACCACAGAAGAAC CCTCTCCAC CCAAGCGGCC GAGAAACCTG CGTGCAATCC GCACCACAGAAGAAC CCTCTCCAC CAACCAGCTT TTGATCAGATC TTGATCCCTT 3720 CTTACCAGAG GGCGCCCCAG CTGGCAATTC CCGTTCGCTT ACTTTGCAGG GCTTCCCAAC 3840 GTCTAGCTAT CGCCATGTAA GCCCACTGCA AGCTACCTGC TTTCTCTTTT CGCTTGCGT 3900 ACTGGCTTTC TACGTGTTCC GCTTCCTTTA GCAGCCCTTG GCTTCTCTTTG GCTTTCTGCGG 3960 ACTGGCTTTC TACGTGTTCC GCTTCCTTTA GCAGCCCTTG GCTTCCTTTT TTGTTAAATC GCCCCTAGAT ACCTCTTT TTACACCAATA 4080 GCCGAAAATC GCCAAAATCC CTTATAAATC AAAAGAACGTC GCTTGCGCA 4020 AAAAACCGTC TATCAGGGCG ATGGCGGATC AGCTTCACTT TTAACCAATA 4080 GCGTAAGGAG AAAATACCGC ATCAGCGCT CTTCCGCTTC CTCCACTA CCCGCACAGAT 4200 AAAAACCGTC TATCAGGGCG ATCAGGCGCT CTTCCGCTTC CTCGCTCACT TCAAAGGGCG 4200 GCGTAAGGAG AAAATACCGC CGAGCGGTA AGCTTATTGCG GTTGAAATA CCGCACAGAT 4200 CCACAGAATC AGGGGATAAC CGGAGAAGA ACATCCGCTC CTCCCCTCC CTCGCTCACT AAAGGCCGAC AAAAGGCCA 4440	CGTATGCAGC (	CGCCGCATTG	CATCACCCAT	CATCCAMACM	GGGCAGGTAG	CCGGATCAAG	
GCGTGCCTCG TCGGGCACG CTGCGCAAGG AACGCCCGTC GTGGCCAGCC ACGATAGCCG 3480 CGGTGCCTCG TCCTGCAGTT CATTCAGGGC ACCGGACAGG TCGGTCTTGA CAAAAAGAAC 3540 TGCCCAGTCA TAGCCGAATA GCCTGCTCAC CCAAGCGGCC GGAGAACCTG CGTGCAATCC 3660 ATCTTGTTCA ATCATGCGAA ACGATCCTCA TCCTGTCTCT TGATCAGATC TTGATCCCTT ACTTGCCAAC GCCGCAATAC CTGGCCATCA ACCATCAGTTT ACTTTGCAGG GCTTCCCAC CTGGCATTC CTGTCTCTT TAATCAGAGC GCTTCCCAC CTGGCAATCC CTGGCAATCC CTGGCAATCC CTGCCATCA AACCGCCCA 3840 GTCTAGCTAT CGCCATGTAA GCCCACTGCA AGTAGCTGC ATCATCAGTTT CTGTTCCTTTT CAGGTGTTC CAGGTAGCCC ACTCAGTTA GCAGCCCTTTA GCAGCACC GCTTCCTTTA GCAGCCCTTTA GCAGCACC GCTTCCTTTA GCAGCCCTTTA GCAGCACC GCTTCCTTTA GCAGCCCTTTA GCAGCCCC AGTAGCACC GCTTCCTTTA GCAGCCCTTTA GCACCAATA ACCCCCCAAAATC GCCCAAAATCC CTTTATAAATC AAAAGAACGTC GCTTCCGCTT TATCACCAATA AAAAAAACCGTC TATCAGGGCG ATGGCGGATC AAAAAAACCGTC TATCAGGGCG ATCAGCGCTT CCGCTCACT TATCAGGGCG AAAAAACCGTC TATCAGGGCG ATCAGCGCTT CCGCCTCACT TACAGGGCG AAAAAACCGCC ATCAGGCGCT CTTCCGCTTC CTCGCTCACT CAAAGGCCGACAGAT ACCGCACAGAT AAAAAACCGTC TCGGCTGC CCACAGATA CCGCACAGAT ACCGCCACAGAT AAAAAACCGTC TCGGCTGC CCACAGATA CCGCACAGAT AAAAAACCGTC TCGGCTGC CCACAGATA CCGCACAGAT AAAAAACCGCC AAAAAAACCGC CCACAGATA CCGCACAGAT AAAAAACCGCC AAAAAACCGC CCACAGATA CCGCACAGAT AAAAAACCGCC AAAAAACCGC CCACAGATA CCGCACAGAT CCACACAGAT CCACACAGAT CCACACAGAT AAAAAACCGC CCACAGATA CCGCACAGAT AAAAAACCGC CCACAGAAAAAACCGC CCACAGAAAAAACCGC CCACAGAAAAAACCGC CCACAGAAAAAACCGC CCACAGAAAAAAACCGC CCACAGAAAAAAACCGC CCACAGAAAAAAACCGC CCACAGAAAAAAACCGC CCACAGAAAAAAAA	AGATGACAGG	AGATCCTGCC	CCCCCACCAC	CCCCLLATIC	TTCTCGGCAG	GAGCAAGGTG	3360
CGGGCGCCCC TGCGCTGACA GCCGGAACAC GGCGCATCA TGCCCAGTCA TAGCCGAATA GCCTCCCCC CCAAGCGGCC ACCAGCCCCA ACCATCCAC CCCAAGCGGCC GGAGAACCT GCGCCATCA ACCATCCAC GCCCCCCCC GCCCCCCCCC GCCCCCCCCC GCCCCCC	AGTGACAACG	TCGAGCACAG	CECCCOTACC	GCCCAATAGC	AGCCAGTCCC	TTCCCGCTTC	3420
CGGGCGCCCC TGCGCTGACA GCCGGAACAC GGCGCATCA TGCCCAGTCA TAGCCGAATA GCCTCCCCC CCAAGCGGCC ACCAGCCCCA ACCATCCAC CCCAAGCGGCC GGAGAACCT GCGCCATCA ACCATCCAC GCCCCCCCC GCCCCCCCCC GCCCCCCCCC GCCCCCC	CGCTGCCTCG	PCCTCCACOO	CIGCOCAAGG	AACGCCCGTC	GTGGCCAGCC	ACGATAGCCG	3480
ATCATGCCAAAAACCGCCAAACCCCCACCCCACCCCACC		CCIGCEGII	CMITCAGGGG	ACCGGACAGG	መር ርርጥ ተመመረ አ	~~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~	3540
ACCTTACCAGA ATCCTTGGGG GCAGAAAGC CATCCAGTTT TGATCAGATC TTGATCCCCT 3720 CTTACCAGAG GGCGCCCAG CTGGCAATC CGGTTCGCTT ACTTTGCAGG GCTTCCCAAC 3780 CTTACCAGAG GGCGCCCAG CTGGCAATC CGGTTCGCTT GCTGTCCATA AAACCGCCCA 3840 TTCCCTTGTC CAGATAGCCC AGTAGCTC ATCATCCGG GGTCAGCACC GTTTCTGTGGG 3960 ACTGGCTTTC TACGTGTTCC GCTTCCTTTA GCAGCCCTTG CGCCCTGAGT GCTTCCGCACAC GCTTCCGCTT TTGTTAAATC AGCTCATTT TTGTTAAATC AGCTCATTT TTAACCAATA 4080 GGCCGAAATC GGCAAAATCC CTTATAAATC ACAAGAACAGT GACTCCAACG TCAAAGGGCG AAAAACCGTC TATCAGGGCG ATGGCGGATC AGCTTATCAG GGTTGAATT TACAGAGAC GCGCACAGAT CGCGCACAGAT AGCTTATCAGGGCG AAAAACCGTC TATCAGGGCG ATCAGGCGCT CTTCCGCTTC CTCGCTCACT TCAAAGGCCG AAAAACCGTC TCGGCTGC CGCCCACGATAG AGCTCCACT TCAAAGGCCG AAAAACCGTC TCGGCTGC CCGCCCACG TCAACGGCGAT 4200 GCCTCGGTCGT TCGGCTGCG CGAGCGGTAT CAGCCTCCACT CTCGCTCACT GACTCGCTGC 4320 CCACAGAATC AGGGGATAAC CCGACAGAT AAAAGGCCAAC CAAAAGGCCAA 4440	TGCCCACTCA	DACCCON NEW	GCCGGAACAC	GGCGGCATCA	GAGCAGCCGA	TTGTCTGTTG	3600
CTTACCAGA GGCGCCCAG CTGGCAATC CGGTTCGCTT ACTTGCAGG GCTTCCCAAC 3840 GTCTAGCTAT CGCCATGTAA GCCCACTGCA AGCTACCTGC TTTCTCTTTG CGCTTGCGTT 3900 ACTGGCTTTC CAGATAGCCC AGTACCC GCTTCCTTTG CGCTTGCGTT 3900 ACTGGCTTTC TACGTGTTCC GCTTCCTTTA ATCACCCG GGTCAGCAC GTTTCTGCGG 3960 GCGTAAAGCT GTCAATTCC GCTTAAATC AGCTCATTT TTAACCAATA 4080 GGCCGAAATC GGCAAAATCC CTTATAAATC AAAAAAATAG CCCGAAGATAG GGTTGAGTGT 4140 AAAAACCGTC TATCAGGGCG ATGGCGGATC AGCTTCAACG TCAAAGGGCG 4200 GCGTAAGGAG AAAATACCGC ATCAGGCGCT CTTCCGCTTC CTCGCTCACT TCAAAGGCCG 4200 GCGTAAGGAG AAAATACCGC ATCAGGCGCT CTTCCGCTTC CTCGCTCACT GACTCCACGT TCAAAGGCCG 4200 GCTCGGTCGT TCGGCTGCG CGAGGGGTAT CAGCTCACCT CTCGCTCC TCGCTCACT AAAGGCCGACAGAT 4320 CCACAGAAATC AGGGGATAAC GCAGGAAAGA ACATGTGAGC AAAAGGCCAG CAAAAGGCCA 4440	ATCTTCTTCT	MCATCOCA TA	GCCTCTCCAC	CCAAGCGGCC	GGAGAACCTG	CGTGCAATCC	3660
GTCTAGCTAT CGCCATGTAA GCCCACTGCA AGCTACCTGC TTTCTCTTTG CGCTTGCGTT 3900 TTCCCTTGTC CAGATAGCCC AGTAGCTGC ATTCATCCGG GGTCAGCAC GTTTCTGCGG 3960 ACTGGCTTTC TACGTGTTCC GCTTCCTTTA GCACCCACTGCA AGCTACCTG GGTCAGCAC GTTTCTGCGG 3960 GCGCGAAATC GCCAATTCC CGTTAAATTT TTGTTAAATC AGCTCATTT TTAACCAATA 4080 GGCCGAAATC GGCAAAATCC CTTATAAATC AAAAGAATAG CCCGAGATAG GGTTGAGTGT 4140 TGTTCCAGTT TGGAACAAGA GTCCACTATT AAAGAACGTG GACTCCAACG TCAAAAGGGCG 4200 GCGTAAGGAG AAAATACCGC ATCAGGCGAT AGCTTATGCG GTGGAAATA CCGCACAGAT 4260 GCGTAAGGAG AAAATACCGC ATCAGGCGCT CTTCCGCTTC CTCGCTCACT GACTCGCTGC 4320 GCTCGGTCGT TCGGCTGCG CGAGGGAAAGA ACATGTGAGC AAAAGGCCAG ATACGGTTAT 4380 CCACAGAAATC AGGGGATAAC GCAGAGAAAAAGGCCA AAAAGGCCAG CAAAAAGGCCA 4440	GCGCCATCAC	ATCATGLGAA .	ACGATCCTCA	TCCTGTCTCT	TGATCAGATC	TTGATCCCCT	3720
TTCCCTTGTC CAGATAGCCC ACCCACGCA AGCTACCTGC TTTCTCTTTG CGCTTGCGTT 3900  TTCCCTTGTC CAGATAGCCC AGTAGCTCC ATTCATCCGG GGTCAGCACC GTTTCTGCGG 3960  ACTGGCTTTC TACGTGTTCC GCTTCCTTTA GCACCCCTTG CGCCCTGAGT GCTTGCGGCA 4020  GCGCGAAATC GGCAAATCCC CGTTAAATTT TTGTTAAATC AGCTCATTT TTAACCAATA 4080  GGCCGAAATC GGCAAAATCC CTTATAAATC AAAAGAATAG CCCGAGATAG GGTTGAGTGT 4140  AAAAACCGTC TATCAGGGCG ATGGCGGATC AGCTTATGCG GACTCCAACG TCAAAGGGCG 4200  GCGTAAGGAG AAAATACCGC ATCAGGGCGT CTTCCGCTTC CTCGCTCACT GACTCGCTGC 4320  GCTCGGTCGT TCGGCTGCG CGAGGGAAAGA ACATGTGAGC AAAAGGCCAG ATACGGTTAT 4380  CCACAGAAATC AGGGGATAAC GCAGGAAAAGA ACATGTGAGC AAAAGGCCAG CAAAAGGCCA 4440	CTTACCACAC	ATCCTTGGCG	GCAAGAAAGC	CATCCAGTTT	ACTTTGCAGG	GCTTCCCAAC	3780
TTCCCTTGTC CAGATAGCCC ACCCACGCA AGCTACCTGC TTTCTCTTTG CGCTTGCGTT 3900  TTCCCTTGTC CAGATAGCCC AGTAGCTCC ATTCATCCGG GGTCAGCACC GTTTCTGCGG 3960  ACTGGCTTTC TACGTGTTCC GCTTCCTTTA GCACCCCTTG CGCCCTGAGT GCTTGCGGCA 4020  GCGCGAAATC GGCAAATCCC CGTTAAATTT TTGTTAAATC AGCTCATTT TTAACCAATA 4080  GGCCGAAATC GGCAAAATCC CTTATAAATC AAAAGAATAG CCCGAGATAG GGTTGAGTGT 4140  AAAAACCGTC TATCAGGGCG ATGGCGGATC AGCTTATGCG GACTCCAACG TCAAAGGGCG 4200  GCGTAAGGAG AAAATACCGC ATCAGGGCGT CTTCCGCTTC CTCGCTCACT GACTCGCTGC 4320  GCTCGGTCGT TCGGCTGCG CGAGGGAAAGA ACATGTGAGC AAAAGGCCAG ATACGGTTAT 4380  CCACAGAAATC AGGGGATAAC GCAGGAAAAGA ACATGTGAGC AAAAGGCCAG CAAAAGGCCA 4440	CTIACCAGAG (	GCGCCCCAG	CTGGCAATTC	CGGTTCGCTT	GCTGTCCATA	AAACCGCCCA	3840
ACTGGCTTC TACGTGTCC GCTTCCTTTA GCAGCCCTG GCTCAGGACC GTTTCTGCGG GCTGGAGACC GCTTGCGTTA GCAGCCCTTG CGCCCTGAGT GCTTGCGGCA 4020 GCGCGAAATC GCCAAATCC CGTTAAATT TTGTTAAATC AGCTCATTT TTAACCAATA 4080 GCCGGAAATC GGCAAAATCC CTTATAAATC AAAAGAATAG CCCGAGAATAG GGTTGAGTGT 4140 AAAAACCGTC TATCAGGGCG ATGGCGGATC AGCTTATGGAATA CCGCACAGGAT 4200 GCGTAAGGAG AAAATACCGC ATCAGGCGCT CTTCCGCTTC CTCGCTCACT GACTCGCTGC 4320 GCTCGGTCGT TCGGCTGCG CGAGGGAAAA ACATCTGAGC AAAAGGCCAG ATACGGTTAT 4380 CCACAGAAATC AGGGGATAAC GCAGGAAAAAAAGGCCA 4440	GICTAGCTAT (	JGCCATGTAA	GCCCACTGCA	AGCTACCTGC	المنتسان المنتسانات	CCCMMCCCMM	
GCGTGAAGCT GTCAATTCCG CGTTAAATTT TTGTTAAATC AGCTCATTT TTAACCAATA 4080 GGCCGAAATC GGCAAAATCC CTTATAAATC AAAAGAATAG CCCGAGATAG GGTTGAGTGT 4140 TGTTCCAGTT TGGAACAAGA GTCCACTATT AAAGAACGTG GACTCCAACG TCAAAGGGCG 4200 AAAAACCGTC TATCAGGGCG ATCGCGGATC AGCTTATGCG GTGGAAATA CCGCACAGAT 4260 GCGTAAGGAG AAAATACCGC ATCAGGCGCT CTTCCGCTTC CTCGCTCACT GACTCGCTGC 4320 GCTCGACAGAATC AGGGGATAAC GCAGGAAAGAAAAGGCCAA AAAAGGCCAG CAAAAGGCCAA 4440	TICCCTTGTC (	AGATAGCCC .	AGTAGCTGAC	ATTCATCCGG	GGTCAGCACC	COMPACTICACA	
GGCCGAAATC GGCAAATCC CTTATAATT TTGTTAAATC AGCTCATTT TTAACCAATA 4080 GGCCGAAATC GGCAAAATCC CTTATAAATC AAAAGAATAG CCCGAGATAG GGTTGAGTGT 4140 TGTTCCAGTT TGGAACAAGA GTCCACTATT AAAGAACGTG GACTCCAACG TCAAAGGGCG 4200 AAAAACCGTC TATCAGGGCG ATCAGGCGCT CTTCCGCTTC CTCGCTCACT GACTCGCTGC 4320 GCTCGGTCGT TCGGCTGCG CGAGCGGTAT CAGCTCACTC AAAGGCCGTA ATACGGTTAT 4380 CCACAGAAATC AGGGGATAAC GCAGGAAAGA ACATCTGAGC AAAAGGCCAG CAAAAGGCCA 4440	weredectate.	PACGIGITICE (	GCTTCCTTTA	GCAGCCCTTG	CGCCCTGAGT	CCMMCCCCCA	
TGTTCCAGTT TGGAACAAGA GTCCACTATT AAAGAACGTG GACTCCAACG TCAAAGGGCG 4200  AAAAACCGTC TATCAGGGCG ATGGCGGATC AGCTTATGCG GTGTGAAATA CCGCACAGAT 4260  GCGTAAGGAG AAAATACCGC ATCAGGCGCT CTTCCGCTTC CTCGCTCACT GACTCGCTGC 4320  GCTCGGTCGT TCGGCTGCG CGAGCGGTAT CAGCTCACTC AAAGGCCGTA ATACGGTTAT 4380  CCACAGAAATC AGGGGATAAC GCAGGAAAGA ACATGTGAGC AAAAGGCCAG CAAAAGGCCA 4440	GCGIGAMGCT (	FICAATTCCG (	CGTTAAATTT	TTGTTAAATC	AGCTCATTT	TTDAACCAAMA	
AAAAACCGTC TATCAGGGCG ATGGCGGATC AGCTTATGCG GACTCCAACG TCAAAGGGCG 4200  AAAAACCGTC TATCAGGGCG ATGGCGGATC AGCTTATGCG GTGTGAAATA CCGCACAGAT 4260  GCGTAAGGAG AAAATACCGC ATCAGGCGCT CTTCCGCTTC CTCGCTCACT GACTCGCTGC 4320  GCTCGGTCGT TCGGCTGCG CGAGCGGTAT CAGCTCACTC AAAGGCCGTA ATACGGTTAT 4380  CCACAGAAATC AGGGGATAAC GCAGGAAAGA ACATGTGAGC AAAAGGCCAG CAAAAGGCCA 4440	GGCCGAAATC (	GCAAAATCC (	CTTATAAATC	AAAAGAATAG	CCCGAGAMAG	CCMMCACMCM	
GCGTAAGAC AAAATACCGC ATCAGGCGATC AGCTTATGCG GTGTGAAATA CCGCACAGAT 4260 GCGTAAGGAG AAAATACCGC ATCAGGCGCT CTTCCGCTTC CTCGCTCACT GACTCGCTGC 4320 GCTCGGTCGT TCGGCTGCG CGAGCGGTAT CAGCTCACTC AAAGGCCGTA ATACGGTTAT 4380 CCACAGAAATC AGGGGATAAC GCAGGAAAGA ACATGTGAGC AAAAGGCCAG CAAAAGGCCA 4440	retrected.	IGGAACAAGA (	GTCCACTATT	AAAGAACGTG	GACTCCAACC	man na adama	
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&2	CCACAGAATC A	AGGGGATAAC	GCAGGAAAGA	ACATGTGAGC	AAAAGGCCAG	CAAAAGGCCA	
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ATCACAAAAA TCGA AGGCGTTTCC CCCT GATACCTGTC CGCC GGTATCTCAG TTCG TTCAGCCCGA CCGC ACGACTTATC GCCA GCGGTGCTAC AGAG TTGGTATCTG CGCTC CCGGCAAACA AACC	GCCGCG TTGCTGGCGT CGCTCA AGTCAGAGGT GGAAGC TCCCTCGTGCA TTTCTC CCTTCGGGAA GTGTAG GTCGTTCGCT TGCGCC TTATCCGGTA CTGGCA GCAGCCACTG TTCTTG AAGTGGTGGC CTGCTG AAGCCAGTTA ACCGCT GGTAGCGGCG CTCAAG AAGATCCTTT	GGCGAAACCC GCTCTCCTGT GCGTGGCGCT CCAAGCTGGG ACTATCGTCT GTAACAGGAT CTAACTACG CCTTCGGAAA	GACAGGACTA TCCGACCCTG TTCTCATAGC CTGTGTGCAC TGAGTCCAAC TAGCAGAGCG CTACACTAGA AAGAGTTGGT	TAAAGATACC CCGCTTACCG TCACGCTGTA GAACCCCCCG CCGGTAAGAC AGGTATGTAG AGGACAGTAT AGCTCTTGAT	4500 4560 4620 4680 4740 4800 4860 4920 4980 5040
CAGAAAAAA GGATO	CTCAAG AAGATCCTTT	GTTTTTTGTT GATCTTTTCT	TGCAAGCAGC TACTGAACGG	AGATTACGCG TGATCCCCAC	5040 5100 5107

# (2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 4818 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)

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(xi)	SEQUENCE	DESCRIPTION	N. SEO ID	NO:4:		
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GAGGTAACTC	CCGTTGCGGT	GCTGTTAACG	GTGGAGGGCA	TTGTATTCTG	ATAAGAGTCA	1500
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CTGGAGAATC	TCCGAGACCT	CCTCCATCTC	CTGGCCTTCT	TGCTGCAGAT	AGCCAATGAC	1980
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CTGAGTAGGT	GTCATTCTAT	TOTOGGGGGGGG	TAATAAAATG	AGGAAATTGC .	ATCGCATTGT	2340
TGGGAAGACA	ATAGCAGGGG	CCTCCCCC *	GGGGTGGGGC	AGGACAGCAA	GGGGAGGAT	2400
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CLICCCCCCLL	CAGTGACAAC	GTCGAGCACA	CCMCCCCC		CAGCCAGTCC CGTGGCCAGC	3120
						3180
						3240
						3300
					CGGAGAACCT	3360
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						3660
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AAGGACAGTA TAGCTCTTGA	TCCGGCAAAC	AAACCACCC	GAAGCCAGTT .	ACCTTCGGAA	Aaagagttgg	4680
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CAGATTACGC (GTGATCCCCA	CCGGAAnn	MUGATUTCAA	GAAGATCCTT	TGATCTTTTC	TTACTGAACG	4800
				•		4818

WHAT IS CLAIMED:

1. A polycationic agent having the following formula:

$$Ta - \begin{bmatrix} R_1 & R_2 & O \\ -N - C - C - C \\ \vdots & R_3 \end{bmatrix} - Tc$$

wherein

n is an integer from 10 to 100;

 R_1 , R_2 , and R_3 for each monomer,

are independently selected from moieties having a molecular weight from 1 to 200 daltons;

Ta and Tc are terminating groups;

R, is not hydrogen for at least one monomer;

wherein said polycationic agent comprises at least 25% positively charged monomers, excluding the terminal groups, and

wherein said polycationic agent exhibits a net positive electrical charge at a physiological pH.

- 2. The polycationic agent according to claim 1, wherein said polycationic agent comprises repeating trimers.
- 3. The polycationic agent according to claim 2, wherein two R_1 groups in each trimer are neutral moieties and one R_1 group in each trimer is a cationic moiety.
- 4. The polycationic agent according to claim 1, wherein R_1 is selected from the group consisting of positively charged, negatively charged and neutral moieties.
- 5. The polycationic agent according to claim 1, wherein R₁ is selected from substituents found on amino acids.

- 6. The polycationic agent of claim 1, wherein R₁ is selected from the group consisting of aromatic and aliphatic groups.
- 7. The polycationic agent according to claim 1, wherein at least one R₁ is selected from the group consisting of alkylammonium, aminoalkyl guanidinoalkyl, amidinoalkyl, piperidyl, guanidinobenzyl, amidinobenzyl, pyridylmethyl, aminobenzyl, alkyphenyl, indolylalkyl, alkoxyphenylalkyl, halophenylalkyl, and hydroxyphenylalkyl.
- 8. The polycationic agent according to claim 3, wherein said cationic moeity is aminoethyl.
- 9. The polycationic agent according to claim 8, wherein said neutral moieties are selected from the group consisting of phenethyl, benzyl, phenylpropyl, (R) alpha-methylbenzyl, (S) alpha-methylbenzyl, methoxyphenethyl, and chlorophenethyl.
- 10. The polycationic agent of claim 1, wherein R_1 and R_3 are both hydrogen for at least one monomer.
 - 11. The polycationic agent of claim 10 wherein n is 36.
 - 12. The polycationic agent of claim 10, wherein n is 24.
 - 13. The polycationic agent of claim 10, wherein n is 18.
 - 14. The polycationic agent of claim 10, wherein n is 12.
- 15. The polycationic agent of claim 8, wherein Ta and Tc are terminal groups selected from the group consisting of polypeptide, lipid, lipoprotein, vitamin, hormone, polyakylene glycol, and saccharide.
 - 16. A composition comprising:

- (i) a polynucleotide; and
- (ii) the polycationic agent of claim 1, wherein said polycationic agent is capable of mediating entry of polynucleotides into a cell.
 - 17. A pharmaceutical composition comprising:
 - (i) a pharmaceutically acceptable carrier;
 - (ii) a therapeutically effective amount of polynucleotides; and
- (iii) an amount effective to neutralize said polynucleotides of the polycationic agent of claim 1 wherein said polycationic agent is capable of mediating entry of polynucleotides into a cell.
- 18. A method of complexing polynucleotides with a polycationic agent comprising:
 - (i) providing a polynucleotide; and
- (ii) contacting said polynucleotide with the polycationic agent of claim 1, wherein said polycationic agent is capable of mediating entry of polynucleotides into a cell.
- 19. A method of condensing polynucleotides, said method comprising: contacting a polynucleotide with a condensing amount of the polycationic agent of claim I, wherein said condensing amount is an amount of polycationic agent sufficient to reduce the size of said polynucleotide.
- 20. A method of inhibiting serum degradation of polynucleotides, said method comprising contacting a polynucleotide with the polycationic agent of claim 1 wherein said polycationic agent is present in an amount effective to inhibit serum degradation by at least 10 minutes.
 - 21. A composition comprising:
 - (i) a lipoprotein;
 - (ii) a polynucleotide binding molecule; and
 - (iii) a polynucleotide,

wherein said composition is capable of increasing the frequency of polynucleotide uptake into a cell.

- 22. The composition of claim 21, wherein the lipoprotein is selected from the group consisting of high density lipoprotein, intermediate density lipoprotein, low density lipoprotein, and very low density lipoprotein,
- 23. The composition of claim 21, wherein the lipoprotein a mutant, fragment or fusion of the protein selected from the group consisting of high density lipoprotein, intermediate density lipoprotein, low density lipoprotein, and very low density lipoprotein.
- 24. The composition of claim 21, wherein the lipoprotein is acetylated low density lipoprotein.
 - 25. A pharmaceutical composition comprising
 - (a) a therapeutically effective amount of a polynucleotide;
- (b) a polynucleotide binding molecule in an amount effective to neutralize the negative charge of said polynucleotide; and
 - (c) a therapeutically effective amount of lipoprotein.
- 26. The pharmaceutical composition of claim 26, wherein said polynucleotide is a polycationic agent.
- 27. A method of producing a composition for facilitating entry of a polynucleotide into a cell said method comprising:
 - (i) providing a polynucleotide
- (ii) providing a polynucleotide binding molecule in an amount effective to neutralize said polynucleotide;
- (ii) contacting said polynucleotide with said polynucleotide binding molecule to form a complex;
 - (iii) providing a lipoprotein; then
 - (iv) contacting the complex with said lipoprotein.

28. A method of increasing the frequency of polynucleotide uptake into a cell said method comprising

- (i) providing a composition that comprises
 - (a) a therapeutically effective amount of a polynucleotide;
- (b) a polynucleotide binding molecule in an amount effective to neutralize said polynucleotide; and
 - (c) an effective amount lipoprotein; then
 - (ii) contacting said composition to said cell.
- 29. A method of increasing the frequency of polynucleotide uptake into a cell said method comprising:
 - (i) providing a composition that comprises
 - (a) a polynucleotide; and
- (b) the polycationic agent of claim 1 in an amount effective to neutralize the negative charge of said polynucleotide; then
 - (ii) contacting said composition to said cell.

Two Step Monomer Assembly

Step 1: Acylation

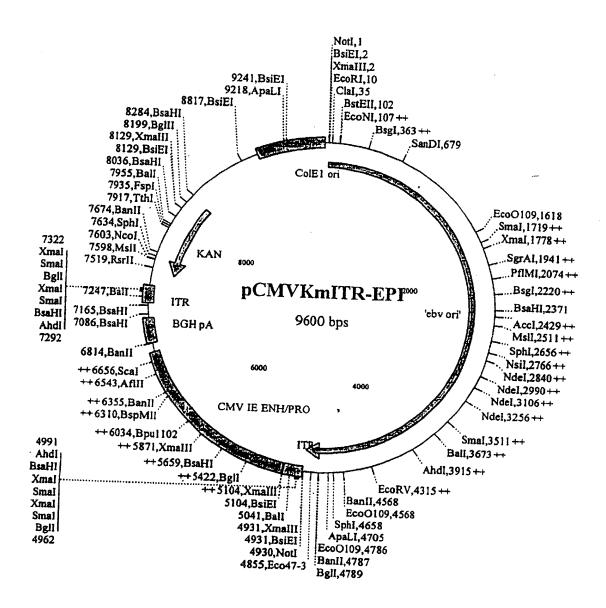
Step 2: Nucleophilic Displacement

Three Step Monomer Assembly

Step 1: Acylation

Step 2: Nucleophilic Displacement

Step 3: Acylation



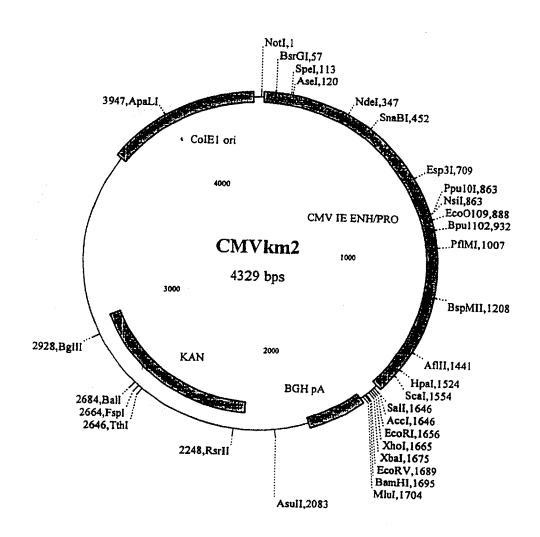
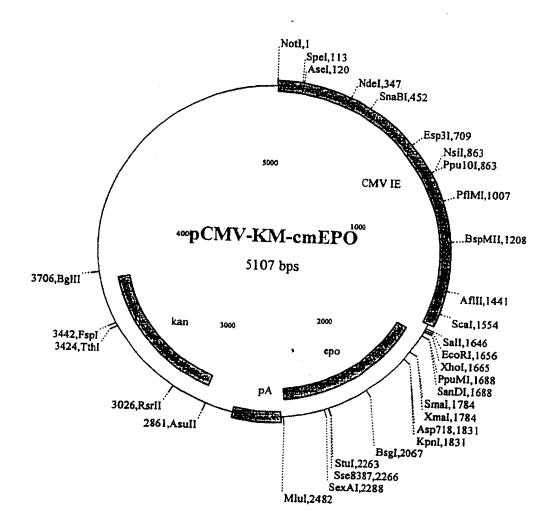
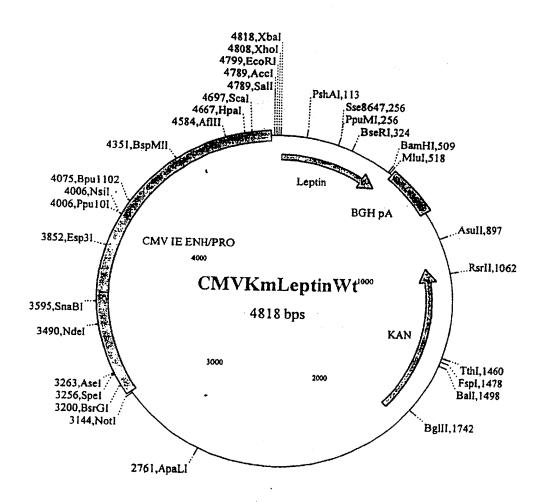
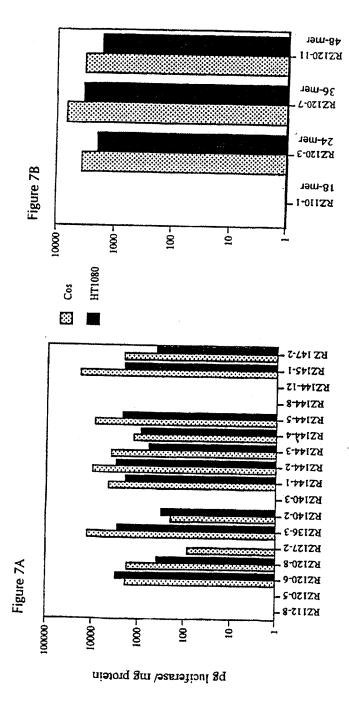


FIGURE 5



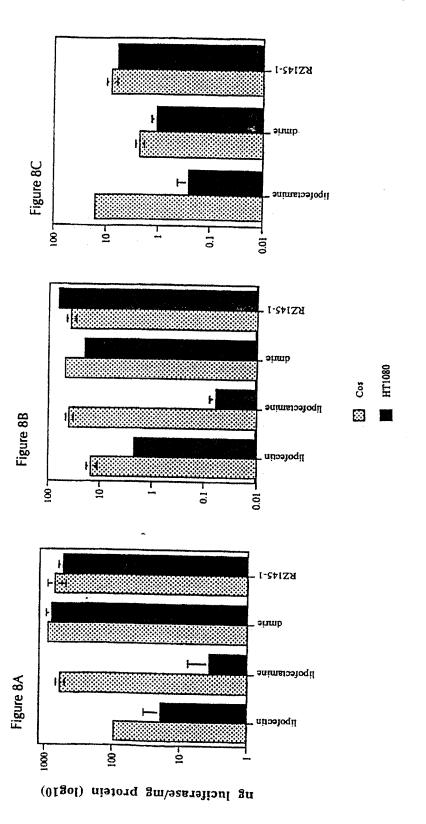


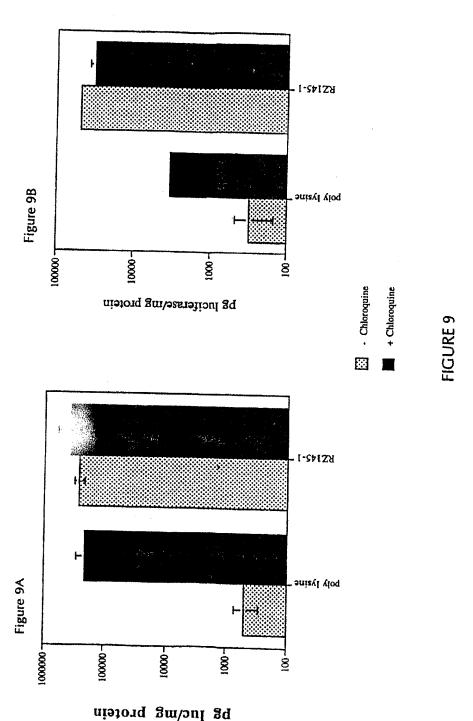




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(30) Priority Data: 60/023,867 13 August 1996 (13.08.96)	ι	Published With international search report. Before the expiration of the time limit for amending the claim and to be republished in the event of the receipt of amendments
(71) Applicant: CHIRON CORPORATION [US/US]; 45 Street, Emeryville, CA 95608 (US).	60 Horto	(88) Date of publication of the international search report: 27 August 1998 (27.08.98
 (72) Inventors: ZUKERMANN, Ronald; 1126 Keele Berkeley, CA 94708 (US). DUBOIS-STRING Nathalie; 1008 The Alameda, Berkeley, CA 94 DWARKI, Varavani; 1177 Old Alameda Pt., Ala 94502 (US). INNIS, Michael, A.; 315 Consta Moraga, CA 94556 (US). MURPHY, John, E.; 49 Court, Oakland, CA 94618 (US). COHEN, Fre Corporation, Intellectual Property - R440, P.O. Emeryville, CA 94662-8097 (US). TETSUO, Warren Drive #530, San Francisco, CA 94131 (U (74) Agents: FUJITA, Sharon, M. et al.; Chiron Co Intellectual Property - R440, P.O. Box 8097, E CA 94662-8097 (US). 	FELLOV 1707 (US ameda, C nce Plac Hourbon eds Chirc Box 809 Uno; 48 JS).	A c, dd n c, do n c, d

(57) Abstract

This invention relates to compositions and methods for increasing the uptake of polynucleotides into cells. Specifically, the invention relates to vectors, targeting ligands, and polycationic agents. The polycationic agents are capable of (1) increasing the frequency of uptake of polynucleotides into a cell, (2) condensing polynucleotides; and (3) inhibiting serum and/or nuclease degradation of polynucleotides.

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C. DOCUME	ENTS CONSIDERED TO BE RELEVANT		
Category °	Citation of document, with indication, where appropriate, of the rela	evant passages	Relevant to claim No.
Х,Р	US 5 679 559 A (KIM JIN-SEOK ET October 1997 see column 2, line 1-20; figure see column 3, line 55-68; claims 1-3,6-8,17; examples 1,3,5,8		1,4,5, 10-29
Х	WO 94 06451 A (CHIRON CORP) 31 M cited in the application see page 34, line 10-20 see page 18; claims 12-22; examp tables 1,5 see page 25, line 22-33		1-29
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<u> </u>	ner documents are listed in the continuation of box C.	Patent family members are listed in anne	ox.
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	NL - 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 651 epo nl, Fax: (+31-70) 340-3016	Gonzalez Ramon, N	

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Box I	Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)
This Inte	emational Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:
1. X	Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:
	Although claims 16,17,21-29 are directed to a method of treatment of the human/animal body, the search has been carried out and based on the alleged effects of the compound/composition.
2. X	Claims Nos.: 1-4 because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
	see FURTHER INFORMATION sheet PCT/ISA/210.
3.	Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).
Box II	Observations where unity of invention is lacking (Continuation of item 2 of first sheet)
This inte	emational Searching Authority found multiple inventions in this international application, as follows:
	A STATE APPROACH, as IOHOWS.
1.	As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2.	As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3.	As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4.	No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:
Remark	con Protest The additional search fees were accompanied by the applicant's protest.
	The daditional solution less were accompanied by the applicant's protest.
	No protest accompanied the payment of additional search fees.

Form PCT/ISA/210 (continuation of first sheet (1)) (July 1992)

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

Claims Nos.: 1-4

In view of the large number of compounds, which are defined by the general definition in the independent claims, the search had to be restricted for economic reasons. The search was limited to the compounds for which pharmacological data was given and/or the compounds mentioned in the claims, and to the general idea underlying the application. (see Guidelines, Chapter III, paragraph 2.3).

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